

SEQUENCE LISTING

<110> WILDT, Stefan MIELE, Robert G. NETT, Juergen H. DAVIDSON, Robert C.

<120> METHODS TO ENGINEER MAMMALIAN-TYPE CARBOHYDRATE STRUCTURES

<130> GF0022P

<140> 10/500,240

<141> 2005-03-23

<150> PCT/US02/41510

<151> 2002-12-24

<150> 60/344,169

<151> 2001-12-27

<160> 106

<170> FastSEQ for Windows Version 4.0

<210> 1

<211> 35

<212> DNA

<213> Artificial Sequence

<220>

<223> Primer

<400> 1

ggtgttttgt tttctagatc tttgcaytay cartt

<210> 2

<211> 36

<212> DNA

<213> Artificial Sequence

<220>

<223> Primer

<400> 2

agaatttggt gggtaagaat tccarcacca ytcrtg

36

35

<210> 3

<211> 32

<212> DNA

<213> Artificial Sequence

<220>

<223> Primer

<400> 3

cctaagctgg tatgcgttct ctttgccata tc	32
<210> 4 <211> 30 <212> DNA <213> Artificial Sequence	
<220> <223> Primer	
<400> 4 gcggcataaa caataataga tgctataaag	30
<210> 5 <211> 20 <212> DNA <213> Artificial Sequence	
<220> <223> Primer	
<400> 5 aattaaccct cactaaaggg	20
<210> 6 <211> 22 <212> DNA <213> Artificial Sequence	
<220> <223> Primer	
<400> 6 gtaatacgac tcactatagg gc	22
<210> 7 <211> 24 <212> DNA <213> Artificial Sequence	
<220>	
<400> 7 ccacatcatc cgtgctacat atag	24
<210> 8 <211> 44 <212> DNA <213> Artificial Sequence	
<220> <223> Primer	
<400> 8 acgaggcaag ctaaacagat ctcgaagtat cgagggttat ccag	44
<210> 9	

<211> 44 <212> DNA <213> Artificial Sequence	
<220> <223> Primer	
<400> 9 ccatccagtg tcgaaaacga gccaatggtt catgtctata aatc	4 4
<210> 10 <211> 24 <212> DNA <213> Artificial Sequence	
<220> <223> Primer	
<400> 10 agcctcagcg ccaacaagcg atgg	24
<210> 11 <211> 44 <212> DNA <213> Artificial Sequence	
<220> <223> Primer	
<400> 11 ctggataacc ctcgatactt cgagatctgt ttagcttgcc tcgt	44
<210> 12 <211> 44 <212> DNA <213> Artificial Sequence	
<220>	
<400> 12 gatttataga catgaaccat tggctcgttt tcgacactgg atgg	44
<210> 13 <211> 20 <212> DNA <213> Artificial Sequence	
<220> <223> Primer	
<400> 13 atcctttacc gatgctgtat	20
<210> 14 <211> 27 <212> DNA <213> Artificial Sequence	

<220> <223> Primer	
<400> 14 ataacagtat gtgttacacg cgtgtag	27
<210> 15 <211> 36 <212> DNA <213> Artificial Sequence	
<220> <223> Primer	
<400> 15 teetggegeg cetteeegag agaactggee teeete	36
<210> 16 <211> 37 <212> DNA <213> Artificial Sequence	
<220> <223> Primer	
<400> 16 aattaattaa ccctagccct ccgctgtatc caacttg	37
<210> 17 <211> 28 <212> DNA <213> Artificial Sequence	
<220> <223> Primer	
<400> 17 aatgagatga ggctccgcaa tggaactg .	28
<210> 18 <211> 28 <212> DNA <213> Artificial Sequence	
<220> <223> Primer	
<400> 18 ctgattgctt atcaacgaga attccttg	28
<210> 19 <211> 28 <212> DNA <213> Artificial Sequence	
<220> <223> Primer	

	<400> tgttg		28
	<210> <211> <212> <213>	29	
	<220> <223>	Primer	
	<400> agaga		29
	<210> <211> <212> <213>	27	
	<220> <223>	Primer	
	<400> aaatc		27
•	<210> <211> <212> <213>	28	
	<220> <223>	Primer	
	<400> agcgat		28
<	<210> <211> <212> <213>	4	
	<400> His As 1	23 sp Glu Leu	
<	<210><211><211><212><213>	458	
<	(222>	MOD_RES (304)(318) Xaa is a variable amino acid	
<	(220>		

<221> MOD RES $\langle 222 \rangle (41\overline{6}) \dots (436)$ <223> Xaa is a variable amino acid <400> 24 Met Glu Gly Glu Gln Ser Pro Gln Gly Glu Lys Ser Leu Gln Arg Lys 10 Gln Phe Val Arg Pro Pro Leu Asp Leu Trp Gln Asp Leu Lys Asp Gly 30 20 25 Val Arg Tyr Val Ile Phe Asp Cys Arg Ala Asn Leu Ile Val Met Pro 40 Leu Leu Ile Leu Phe Glu Ser Met Leu Cys Lys Ile Ile Ile Lys Lys Val Ala Tyr Thr Glu Ile Asp Tyr Lys Ala Tyr Met Glu Gln Ile Glu Met Ile Gln Leu Asp Gly Met Leu Asp Tyr Ser Gln Val Ser Gly Gly 90 Thr Gly Pro Leu Val Tyr Pro Ala Gly His Val Leu Ile Tyr Lys Met 105 110 Met Tyr Trp Leu Thr Glu Gly Met Asp His Val Glu Arg Gly Gln Val 120 125 Phe Phe Arg Tyr Leu Tyr Leu Leu Thr Leu Ala Leu Gln Met Ala Cys 135 Tyr Tyr Leu Leu His Leu Pro Pro Trp Cys Val Val Leu Ala Cys Leu 150 155 Ser Lys Arg Leu His Ser Ile Tyr Val Leu Arg Leu Phe Asn Asp Cys 165 170 175 Phe Thr Thr Leu Phe Met Val Val Thr Val Leu Gly Ala Ile Val Ala 180 185 Ser Arg Cys His Gln Arg Pro Lys Leu Lys Lys Ser Leu Ala Leu Val 195 200 Ile Ser Ala Thr Tyr Ser Met Ala Val Ser Ile Lys Met Asn Ala Leu 215 Leu Tyr Phe Pro Ala Met Met Ile Ser Leu Phe Ile Leu Asn Asp Ala 230 235 Asn Val Ile Leu Thr Leu Leu Asp Leu Val Ala Met Ile Ala Trp Gln 250 Val Ala Val Ala Val Pro Phe Leu Arg Ser Phe Pro Gln Gln Tyr Leu 265 His Cys Ala Phe Asn Phe Gly Arg Lys Phe Met Tyr Gln Trp Ser Ile 280 285 Asn Trp Gln Met Met Asp Glu Glu Ala Phe Asn Asp Lys Arg Phe Xaa 295 300 310 315 Thr Arg Tyr Pro Arg Ile Leu Pro Asp Leu Trp Ser Ser Leu Cys His 325 330 335 Pro Leu Arg Lys Asn Ala Val Leu Asn Ala Asn Pro Ala Lys Thr Ile 340 345 Pro Phe Val Leu Ile Ala Ser Asn Phe Ile Gly Val Leu Phe Ser Arg 360 Ser Leu His Tyr Gln Phe Leu Ser Trp Tyr His Trp Thr Leu Pro Ile 375 380 Leu Ile Phe Trp Ser Gly Met Pro Phe Phe Val Gly Pro Ile Trp Tyr 390 395 Val Leu His Glu Trp Cys Trp Asn Ser Tyr Pro Pro Asn Ser Gln Xaa 405 410

```
420
                              425
Xaa Xaa Xaa Ser Gly Ser Val Ala Leu Ala Lys Ser His Leu Arg
                           440
Thr Thr Ser Ser Met Glu Lys Lys Leu Asn
                       455
<210> 25
<211> 458
<212> PRT
<213> Saccharomyces cerevisiae
<400> 25
Met Glu Gly Glu Gln Ser Pro Gln Gly Glu Lys Ser Leu Gln Arg Lys
Gln Phe Val Arg Pro Pro Leu Asp Leu Trp Gln Asp Leu Lys Asp Gly
                               25
Val Arg Tyr Val Ile Phe Asp Cys Arg Ala Asn Leu Ile Val Met Pro
                          40
Leu Leu Ile Leu Phe Glu Ser Met Leu Cys Lys Ile Ile Lys Lys
                      55
Val Ala Tyr Thr Glu Ile Asp Tyr Lys Ala Tyr Met Glu Gln Ile Glu
                  70
                                     75
Met Ile Gln Leu Asp Gly Met Leu Asp Tyr Ser Gln Val Ser Gly Gly
                                 90
Thr Gly Pro Leu Val Tyr Pro Ala Gly His Val Leu Ile Tyr Lys Met
                              105
                                                 110
Met Tyr Trp Leu Thr Glu Gly Met Asp His Val Glu Arg Gly Gln Val
115
                         120
Phe Phe Arg Tyr Leu Tyr Leu Leu Thr Leu Ala Leu Gln Met Ala Cys
                      135
Tyr Tyr Leu Leu His Leu Pro Pro Trp Cys Val Val Leu Ala Cys Leu
                                      155
Ser Lys Arg Leu His Ser Ile Tyr Val Leu Arg Leu Phe Asn Asp Cys
                                  170
Phe Thr Thr Leu Phe Met Val Val Thr Val Leu Gly Ala Ile Val Ala
                              185
Ser Arg Cys His Gln Arg Pro Lys Leu Lys Lys Ser Leu Ala Leu Val
                          200
                                              205
Ile Ser Ala Thr Tyr Ser Met Ala Val Ser Ile Lys Met Asn Ala Leu
                       215
                                          220
Leu Tyr Phe Pro Ala Met Met Ile Ser Leu Phe Ile Leu Asn Asp Ala
                  230
                                      235
Asn Val Ile Leu Thr Leu Leu Asp Leu Val Ala Met Ile Ala Trp Gln
                                  250
              245
Val Ala Val Ala Val Pro Phe Leu Arg Ser Phe Pro Gln Gln Tyr Leu
           260
                              265
His Cys Ala Phe Asn Phe Gly Arg Lys Phe Met Tyr Gln Trp Ser Ile
                          280
                                              285
Asn Trp Gln Met Met Asp Glu Glu Ala Phe Asn Asp Lys Arg Phe His
                      295
                                          300
Leu Ala Leu Leu Ile Ser His Leu Ile Ala Leu Thr Thr Leu Phe Val
                   310
                                      315
Thr Arg Tyr Pro Arg Ile Leu Pro Asp Leu Trp Ser Ser Leu Cys His
                           . 330
              325
Pro Leu Arg Lys Asn Ala Val Leu Asn Ala Asn Pro Ala Lys Thr Ile
                              345
Pro Phe Val Leu Ile Ala Ser Asn Phe Ile Gly Val Leu Phe Ser Arg
```

```
360
Ser Leu His Tyr Gln Phe Leu Ser Trp Tyr His Trp Thr Leu Pro Ile
                        375
                                            380
Leu Ile Phe Trp Ser Gly Met Pro Phe Phe Val Gly Pro Ile Trp Tyr
                    390
                                        395
Val Leu His Glu Trp Cys Trp Asn Ser Tyr Pro Pro Asn Ser Gln Ala
                405
                                    410
Ser Thr Leu Leu Ala Leu Asn Thr Val Leu Leu Leu Leu Ala
            420
                                425
Leu Thr Gln Leu Ser Gly Ser Val Ala Leu Ala Lys Ser His Leu Arg
                            440
Thr Thr Ser Ser Met Glu Lys Lys Leu Asn
                        455
<210> 26
<211> 443
<212> PRT
<213> Saccharomyces cerevisiae
<220>
<221> MOD RES
<222> (333)...(347)
<223> Xaa is a variable amino acid
<400> 26
Trp Gln Asp Leu Lys Asp Gly Val Arg Tyr Val Ile Phe Asp Cys Arg
                                    10
Ala Asn Leu Ile Val Met Pro Leu Leu Ile Leu Phe Glu Ser Met Leu
                                25
Cys Lys Ile Ile Ile Lys Lys Val Ala Tyr Thr Glu Ile Asp Tyr Lys
                            40
Ala Tyr Met Glu Gln Ile Glu Met Ile Gln Leu Asp Gly Met Leu Asp
Tyr Ser Gln Val Ser Gly Gly Thr Gly Pro Leu Val Tyr Pro Ala Gly
His Val Leu Ile Tyr Lys Met Met Tyr Trp Leu Thr Glu Gly Met Asp
                                    90
His Val Glu Arg Gly Gln Val Phe Phe Arg Tyr Leu Tyr Leu Leu Thr
                               105
Leu Ala Leu Gln Met Ala Cys Tyr Tyr Leu Leu His Leu Pro Pro Trp
                           120
                                                125
Cys Val Val Leu Ala Cys Leu Ser Lys Arg Leu His Ser Ile Tyr Val
                       135
                                            140
Leu Arg Leu Phe Asn Asp Cys Phe Thr Thr Leu Phe Met Val Val Thr
                   150
                                        155
Val Leu Gly Ala Ile Val Ala Ser Arg Cys His Gln Arg Pro Lys Leu
                165
                                    170
Lys Lys His Gln Thr Cys Lys Val Pro Pro Phe Val Phe Phe Met
                               185
Cys Cys Ala Ser Tyr Arg Val His Ser Ile Phe Val Leu Arg Leu Phe
                           200
Asn Asp Pro Val Ala Met Val Leu Leu Phe Leu Ser Ile Asn Leu Leu
                       215
                                            220
Leu Ala Gln Arg Trp Gly Trp Gly Ser Leu Ala Leu Val Ile Ser Ala
                   230
                                    235
Thr Tyr Ser Met Ala Val Ser Ile Lys Met Asn Ala Leu Leu Tyr Phe
               245
                                   250
```

```
Pro Ala Met Met Ile Ser Leu Phe Ile Leu Asn Asp Ala Asn Val Ile
                             265
Leu Thr Leu Leu Asp Leu Val Ala Met Ile Ala Trp Gln Val Ala Val
                         280
Ala Val Pro Phe Leu Arg Ser Phe Pro Gln Gln Tyr Leu His Cys Ala
                      295
                                        300
Phe Asn Phe Gly Arg Lys Phe Met Tyr Gln Trp Ser Ile Asn Trp Gln
                  310
                                     315
Met Met Asp Glu Glu Ala Phe Asn Asp Lys Arg Phe Xaa Xaa Xaa
              325
                                 330
345
Pro Arg Ile Leu Pro Asp Leu Trp Ser Ser Leu Cys His Pro Leu Arg
                         360
Lys Asn Ala Val Leu Asn Ala Asn Pro Ala Lys Thr Ile Pro Phe Val
                      375
                                        380
Leu Ile Ala Ser Asn Phe Ile Gly Val Leu Phe Ser Arg Ser Leu His
                  390
                                     395
Tyr Gln Phe Leu Ser Trp Tyr His Trp Thr Leu Pro Ile Leu Ile Phe
              405
                                 410
Trp Ser Gly Met Pro Phe Phe Val Gly Pro Ile Trp Tyr Val Leu His
                             425
Glu Trp Cys Trp Asn Ser Tyr Pro Pro Asn Ser
                         440
```

<210> 27 <211> 373

<212> PRT

<213> Homo sapiens

<400> 27

Trp Gln Glu Arg Arg Leu Leu Arg Glu Pro Arg Tyr Thr Leu Leu Val Ala Ala Cys Leu Cys Leu Ala Glu Val Gly Ile Thr Phe Trp Val 25 Ile His Arg Val Ala Tyr Thr Glu Ile Asp Trp Lys Ala Tyr Met Ala 40 45 Glu Val Glu Gly Val Gly Thr Tyr Asp Tyr Thr Gln Leu Gln Gly Asp 55 Thr Gly Pro Leu Val Tyr Pro Ala Gly Phe Val Tyr Ile Phe Met Gly 70 75 Leu Tyr Tyr Ala Thr Ser Arg Gly Thr Asp Ile Arg Met Ala Gln Asn 8.5 90 Ile Phe Ala Val Leu Tyr Leu Ala Thr Leu Leu Leu Val Phe Leu Ile 100 105 Tyr His Gln Thr Cys Lys Val Pro Pro Phe Val Phe Phe Met Cys 120 Cys Ala Ser Tyr Arg Val His Ser Ile Phe Val Leu Arg Leu Phe Asn 135 140 Asp Pro Val Ala Met Val Leu Leu Phe Leu Ser Ile Asn Leu Leu 150 155 Ala Gln Arg Trp Gly Trp Gly Cys Cys Phe Phe Ser Leu Ala Val Ser 165 170 Val Lys Met Asn Val Leu Leu Phe Ala Pro Gly Leu Leu Phe Leu Leu 180 185. Leu Thr Gln Phe Gly Phe Arg Gly Ala Leu Pro Lys Leu Gly Ile Cys 195 200

```
Ala Gly Leu Gln Val Val Leu Gly Leu Pro Phe Leu Leu Glu Asn Pro
                        215
Ser Gly Tyr Leu Ser Arg Ser Phe Asp Leu Gly Arg Gln Phe Leu Phe
                    230
                                        235
His Trp Thr Val Asn Trp Arg Phe Leu Pro Glu Ala Leu Phe Leu His
                245
                                    250
Arg Ala Phe His Leu Ala Leu Leu Thr Ala His Leu Thr Leu Leu Leu
                                265
Leu Phe Ala Leu Cys Arg Trp His Arg Thr Gly Glu Ser Ile Leu Ser
                            280
Leu Leu Arg Asp Pro Ser Lys Arg Lys Val Pro Pro Gln Pro Leu Thr
                        295
Pro Asn Gln Ile Val Ser Thr Leu Phe Thr Ser Asn Phe Ile Gly Ile
                                        315
Cys Phe Ser Arg Ser Leu His Tyr Gln Phe Tyr Val Trp Tyr Phe His
                325
                                    330
Thr Leu Pro Tyr Leu Leu Trp Ala Met Pro Ala Arg Trp Leu Thr His
                                345
Leu Leu Arg Leu Leu Val Leu Gly Leu Ile Glu Leu Ser Trp Asn Thr
       355
                           360
Tyr Pro Ser Thr Ser
   370
```

<210> 28

<211> 269

<212> PRT

<213> Saccharomyces cerevisiae

<400> 28

Val Arg Tyr Val Ile Phe Asp Cys Arg Ala Asn Leu Ile Val Met Pro Leu Leu Ile Leu Phe Glu Ser Met Leu Cys Lys Ile Ile Ile Lys Lys Val Ala Tyr Thr Glu Ile Asp Tyr Lys Ala Tyr Met Glu Gln Ile Glu Met Ile Gln Leu Asp Gly Met Leu Asp Tyr Ser Gln Val Ser Gly Gly Thr Gly Pro Leu Val Tyr Pro Ala Gly His Val Leu Ile Tyr Lys Met 7.5 Met Tyr Trp Leu Thr Glu Gly Met Asp His Val Glu Arg Gly Gln Val 90 Phe Phe Arg Tyr Leu Tyr Leu Leu Thr Leu Ala Leu Gln Met Ala Cys 100 105 Tyr Tyr Leu Leu His Pro Trp Cys Val Val Leu Ala Cys Leu Ser Lys 120 Arg Leu His Ser Ile Tyr Val Leu Arg Leu Phe Asn Asp Cys Phe Thr 135 Thr Leu Phe Met Val Val Thr Val Leu Gly Ala Ile Val Ala Ser Arg 150 155 Cys His Gln Arg Pro Lys Leu Lys Lys Ser Leu Ala Leu Val Ile Ser 170 Ala Thr Tyr Ser Met Ala Val Ser Ile Lys Met Asn Ala Leu Leu Tyr 185 Phe Pro Ala Met Met Ile Ser Leu Phe Ile Leu Asn Asp Ala Asn Val 205 200 Ile Leu Thr Leu Leu Asp Leu Val Ala Met Ile Ala Trp Gln Val Ala 215

```
230
                                         235
Ala Phe Asn Phe Gly Arg Lys Phe Met Tyr Gln Trp Ser Ile Asn Trp
                245
                                    250
Gln Met Met Asp Glu Glu Ala Phe Asn Asp Lys Arg Phe
            2.60
                                265
<210> 29
<211> 258
<212> PRT
<213> Drosophila virilis
<400> 29
Ile Lys Tyr Leu Ala Phe Glu Pro Ala Ala Leu Pro Ile Val Ser Val
                                    10
Leu Ile Val Leu Ala Glu Ala Val Ile Asn Val Leu Val Ile Gln Arg
Val Pro Tyr Thr Glu Ile Asp Trp Lys Ala Tyr Met Gln Glu Cys Glu
                            40
Gly Phe Leu Asn Gly Thr Thr Asn Tyr Ser Leu Leu Arg Gly Asp Thr
                        55
                                            60
Gly Pro Leu Val Tyr Pro Ala Ala Phe Val Tyr Ile Tyr Ser Gly Leu
                    70
                                        75
Tyr Tyr Leu Thr Gly Gln Gly Thr Asn Val Arg Leu Ala Gln Tyr Ile
               85
                                    90
Phe Ala Cys Ile Tyr Leu Leu Gln Met Cys Leu Val Leu Arg Leu Tyr
                                105
Thr Lys Ser Arg Lys Val Pro Pro Tyr Val Leu Val Leu Ser Ala Phe
                            120
Thr Ser Tyr Arg Ile His Ser Ile Tyr Val Leu Arg Leu Phe Asn Asp
                       135
Pro Val Ala Ile Leu Leu Leu Tyr Ala Ala Leu Asn Leu Phe Leu Asp
                   150
                                        155
Gln Arg Trp Thr Leu Gly Ser Ile Cys Tyr Ser Leu Ala Val Gly Val
                                    170
Lys Met Asn Ile Leu Leu Phe Ala Pro Ala Leu Leu Phe Tyr Leu
                               185
                                                    190
Ala Asn Leu Gly Val Leu Arg Thr Leu Val Gln Leu Thr Ile Cys Ala
                            200
                                                205
Val Leu Gln Leu Phe Ile Gly Ala Pro Phe Leu Arg Thr His Pro Met
                       215
                                           220
Glu Tyr Leu Arg Gly Ser Phe Asp Leu Gly Arg Ile Phe Glu His Lys
                   230
                                        235
Trp Thr Val Asn Tyr Arg Phe Leu Ser Lys Glu Leu Phe Glu Gln Arg
                245
                                    250
Glu Phe
<210> 30
<211> 267
<212> PRT
<213> Saccharomyces cerevisiae
Arg Tyr Val Ile Phe Asp Cys Arg Ala Asn Leu Ile Val Met Pro Leu
```

Val Ala Val Pro Phe Leu Arg Ser Phe Pro Gln Gln Tyr Leu His Cys

```
Leu Ile Leu Phe Glu Ser Met Leu Cys Lys Ile Ile Ile Lys Lys Val
                                25
Ala Tyr Thr Glu Ile Asp Tyr Lys Ala Tyr Met Glu Gln Ile Glu Met
Ile Gln Leu Asp Gly Met Leu Asp Tyr Ser Gln Val Ser Gly Gly Thr
Gly Pro Leu Val Tyr Pro Ala Gly His Val Leu Ile Tyr Lys Met Met
Tyr Trp Leu Thr Glu Gly Met Asp His Val Glu Arg Gly Gln Val Phe
Phe Arg Tyr Leu Tyr Leu Leu Thr Leu Ala Leu Gln Met Ala Cys Tyr
                                105
Tyr Leu Leu His Trp Cys Val Val Leu Ala Cys Leu Ser Lys Arg Leu
                            120
His Ser Ile Tyr Val Leu Arg Leu Phe Asn Asp Cys Phe Thr Thr Leu
                        135
                                            140
Phe Met Val Val Thr Val Leu Gly Ala Ile Val Ala Ser Arg Cys His
                   150
                                        155
Gln Arg Pro Lys Leu Lys Lys Ser Leu Ala Leu Val Ile Ser Ala Thr
                                   170
               165
Tyr Ser Met Ala Val Ser Ile Lys Met Asn Ala Leu Leu Tyr Phe Pro
           180
                               185
                                                   190
Ala Met Met Ile Ser Leu Phe Ile Leu Asn Asp Ala Asn Val Ile Leu
                           200
                                                205
Thr Leu Leu Asp Leu Val Ala Met Ile Ala Trp Gln Val Ala Val Ala
                       215
                                           220
Val Pro Phe Leu Arg Ser Phe Pro Gln Gln Tyr Leu His Cys Ala Phe
                   230
                                       235
Asn Phe Gly Arg Lys Phe Met Tyr Gln Trp Ser Ile Asn Trp Gln Met
              245
Met Asp Glu Glu Ala Phe Asn Asp Lys Arg Phe
```

```
<210> 31 <211> 257
```

<212> PRT

<213> Drosophila melanogaster

<400> 31

Lys Tyr Leu Leu Glu Pro Ala Ala Leu Pro Ile Val Gly Leu Phe 10 Val Leu Leu Ala Glu Leu Val Ile Asn Val Val Ile Gln Arg Val 20 Pro Tyr Thr Glu Ile Asp Trp Val Ala Tyr Met Gln Glu Cys Glu Gly 40 Phe Leu Asn Gly Thr Thr Asn Tyr Ser Leu Leu Arg Gly Asp Thr Gly Pro Leu Val Tyr Pro Ala Ala Phe Val Tyr Ile Tyr Ser Ala Leu Tyr 75 Tyr Val Thr Ser His Gly Thr Asn Val Arg Leu Ala Gln Tyr Ile Phe 90 Ala Gly Ile Tyr Leu Leu Gln Leu Ala Leu Val Leu Arg Leu Tyr Ser 105 Lys Ser Arg Lys Val Pro Pro Tyr Val Leu Val Leu Ser Ala Phe Thr 120 125 Ser Tyr Arg Ile His Ser Ile Tyr Val Leu Arg Leu Phe Asn Asp Pro 135

```
Val Ala Val Leu Leu Tyr Ala Ala Leu Asn Leu Phe Leu Asp Arg
                    150
                                         155
Arg Trp Thr Leu Gly Ser Thr Phe Phe Ser Leu Ala Val Gly Val Lys
                165
                                     170
Met Asn Ile Leu Leu Phe Ala Pro Ala Leu Leu Phe Tyr Leu Ala
                                 185
                                                     190
Asn Leu Gly Leu Leu Arg Thr Ile Leu Gln Leu Ala Val Cys Gly Val
                             200
Ile Gln Leu Leu Gly Ala Pro Phe Leu Leu Thr His Pro Val Glu
                         215
Tyr Leu Arg Gly Ser Phe Asp Leu Gly Arg Ile Phe Glu His Lys Trp
225
                    230
                                        235
Thr Val Asn Tyr Arg Phe Leu Ser Arg Asp Val Phe Glu Asn Arg Thr
                245
                                     250
Phe
<210> 32
<211> 1377
<212> DNA
<213> Saccharomyces cerevisiae
<400> 32
atggaaggtg aacagtctcc gcaaggtgaa aagtctctgc aaaggaagca atttgtcaga 60
cctccgctgg atctgtggca ggatctcaag gacggtgtgc gctacgtgat cttcgattgt 120
agggccaatc ttatcgttat gccccttttg attttgttcg aaagcatgct gtgcaagatt 180
atcattaaga aggtagetta cacagagate gattacaagg egtacatgga geagategag 240
atgattcagc tcgatggcat gctggactac tctcaggtga gtggtggaac gggcccgctg 300
gtgtatccag caggccacgt cttgatctac aagatgatgt actggctaac agagggaatg 360
gaccacgttg agcgcgggca agtgtttttc agatacttgt atctccttac actggcgtta 420
caaatggcgt gttactacct tttacatcta ccaccgtggt gtgtggtctt ggcgtgcctc 480
tctaaaagat tgcactctat ttacgtgcta cggttattca atgattgctt cactactttg 540
tttatggtcg teacggtttt gggggctatc gtggccagca ggtgccatca gcgccccaaa 600
ttaaagaagt cccttgcgct ggtgatctcc gcaacataca gtatggctgt gagcattaag 660
atgaatgcgc tgttgtattt ccctgcaatg atgatttctc tattcatcct taatgacgcg 720
aacgtaatcc ttactttgtt ggatctcgtt gcgatgattg catggcaagt cgcagttgca 780
gtgcccttcc tgcgcagctt tccgcaacag tacctgcatt gcgcttttaa tttcggcagg 840
aagtttatgt accaatggag tatcaattgg caaatgatgg atgaagaggc tttcaatgat 900
aagaggttcc acttggccct tttaatcagc cacctgatag cgctcaccac actgttcgtc 960
acaagatacc ctcgcatcct gcccgattta tggtcttccc tgtgccatcc gctgaggaaa 1020
aatgcagtgc tcaatgccaa tcccgccaag actattccat tcgttctaat cgcatccaac 1080
ttcatcggcg tcctattttc aaggtccctc cactaccagt ttctatcctg gtatcactgg 1140
actttgccta tactgatctt ttggtcggga atgcccttct tcgttggtcc catttggtac 1200
gtcttgcacg agtggtgctg gaattcctat ccaccaaact cacaagcaag cacgctattg 1260
ttggcattga atactgttct gttgcttcta ttggccttga cgcagctatc tggttcggtc 1320
gccctcgcca aaagccatct tcgtaccacc agctctatgg aaaaaaagct caactga
<210> 33
<211> 458
<212> PRT
<213> Saccharomyces cerevisiae
Met Glu Gly Glu Gln Ser Pro Gln Gly Glu Lys Ser Leu Gln Arg Lys
                                    10
Gln Phe Val Arg Pro Pro Leu Asp Leu Trp Gln Asp Leu Lys Asp Gly
```

25

30

20

```
Val Arg Tyr Val Ile Phe Asp Cys Arg Ala Asn Leu Ile Val Met Pro
Leu Leu Ile Leu Phe Glu Ser Met Leu Cys Lys Ile Ile Lys Lys
Val Ala Tyr Thr Glu Ile Asp Tyr Lys Ala Tyr Met Glu Gln Ile Glu
                   70
                                       75
Met Ile Gln Leu Asp Gly Met Leu Asp Tyr Ser Gln Val Ser Gly Gly
               85
                                   90
Thr Gly Pro Leu Val Tyr Pro Ala Gly His Val Leu Ile Tyr Lys Met
           100
                               105
Met Tyr Trp Leu Thr Glu Gly Met Asp His Val Glu Arg Gly Gln Val
                           120
Phe Phe Arg Tyr Leu Tyr Leu Leu Thr Leu Ala Leu Gln Met Ala Cys
                       135
Tyr Tyr Leu Leu His Leu Pro Pro Trp Cys Val Val Leu Ala Cys Leu
                   150
                                       155
Ser Lys Arg Leu His Ser Ile Tyr Val Leu Arg Leu Phe Asn Asp Cys
               165
                                   170
                                                       175
Phe Thr Thr Leu Phe Met Val Val Thr Val Leu Gly Ala Ile Val Ala
           180
                               185
Ser Arg Cys His Gln Arg Pro Lys Leu Lys Lys Ser Leu Ala Leu Val
                           200
       195
                                               205
Ile Ser Ala Thr Tyr Ser Met Ala Val Ser Ile Lys Met Asn Ala Leu
                       215
                                          220
Leu Tyr Phe Pro Ala Met Met Ile Ser Leu Phe Ile Leu Asn Asp Ala
                   230
                                       235
Asn Val Ile Leu Thr Leu Leu Asp Leu Val Ala Met Ile Ala Trp Gln
                                   250
              245
Val Ala Val Ala Val Pro Phe Leu Arg Ser Phe Pro Gln Gln Tyr Leu
                              265
His Cys Ala Phe Asn Phe Gly Arg Lys Phe Met Tyr Gln Trp Ser Ile
                          280
Asn Trp Gln Met Met Asp Glu Glu Ala Phe Asn Asp Lys Arg Phe His
                      295
Leu Ala Leu Leu Ile Ser His Leu Ile Ala Leu Thr Thr Leu Phe Val
                   310
                                       315
Thr Arg Tyr Pro Arg Ile Leu Pro Asp Leu Trp Ser Ser Leu Cys His
               325
                                   330
Pro Leu Arg Lys Asn Ala Val Leu Asn Ala Asn Pro Ala Lys Thr Ile
                               345
                                                  350
Pro Phe Val Leu Ile Ala Ser Asn Phe Ile Gly Val Leu Phe Ser Arg
                           360
                                               365
Ser Leu His Tyr Gln Phe Leu Ser Trp Tyr His Trp Thr Leu Pro Ile
                       375
                                           380
Leu Ile Phe Trp Ser Gly Met Pro Phe Phe Val Gly Pro Ile Trp Tyr
                   390
                                       395
Val Leu His Glu Trp Cys Trp Asn Ser Tyr Pro Pro Asn Ser Gln Ala
               405
                                   410
Ser Thr Leu Leu Ala Leu Asn Thr Val Leu Leu Leu Leu Ala
                               425
Leu Thr Gln Leu Ser Gly Ser Val Ala Leu Ala Lys Ser His Leu Arg
                           440
Thr Thr Ser Ser Met Glu Lys Lys Leu Asn
```

<210> 34 <211> 1395

<212> DNA <213> Pichia pastoris <400> 34 atgcctccga tagagccagc tgaaaggcca aagcttacgc tgaaaaatgt tatcggtgat 60 ctagtggctc ttattcaaaa cgttttattt aacccagatt ttagtgtctt cgttgcacct 120 cttttatggt tagctgattc cattgttatc aaggtgatca ttggcactgt ttcctacaca 180 gatattgatt tttcttcata tatgcaacaa atctttaaaa ttcgacaagg agaattagat 240 tatagcaaca tatttggtga caccggtcca ttggtttacc cagccggcca tgttcatgct 300 tactcagtac tttcgtggta cagtgatggt ggagaagacg tcagtttcgt tcaacaagca 360 tttggttggt tatacctagg ttgcttgtta ctatccatca gctcctactt tttctctggc 420 ttagggaaaa tacctccggt ttattttgtt ttgttggtag cgtccaagag actgcattca 480 atatttgtat tgagactctt caatgactgt ttaacaacat ttttgatgtt ggcaactata 540 atcatectte aacaageaag tagetggagg aaagatggea caactattee attatetgte 600 cctgatgctg cagatacgta cagtttagcc atctctgtaa agatgaatgc gctgctatac 660 ctcccagcat tcctactact catatatctc atttgtgacg aaaatttgat taaagccttg 720 gcacctgttc tagttttgat attggtgcaa gtaggagtcg gttattcgtt cattttaccg 780 ttgcactatg atgatcaggc aaatgaaatt cgttctgcct actttagaca ggcttttgac 840 tttagtcgcc aatttcttta taagtggacg gttaattggc gctttttgag ccaagaaact 900 ttcaacaatg tccattttca ccagctcctg tttgctctcc atattattac gttagtcttg 960 ttcatcctca agttcctctc tcctaaaaac attggaaaac cgcttggtag atttgtgttg 1020 gacattttca aattttggaa gccaacctta tctccaacca atattatcaa cgacccagaa 1080 agaagcccag attttgttta caccgtcatg gctactacca acttaatagg ggtgcttttt 1140 gcaagatett tacactacca gtteetaage tggtatgegt tetetttgee atateteett 1200 tacaaggete gtetgaactt tatageatet attattgttt atgeegetea egagtattge 1260 tggttggttt tcccagctac agaacaaagt tccgcgttgt tggtatctat cttactactt 1320 atcctgattc tcatttttac caacgaacag ttatttcctt ctcaatcggt ccctgcagaa 1380 aaaaagaata cataa 1395 <210> 35 <211> 464 <212> PRT <213> Pichia pastoris <400> 35 Met Pro Pro Ile Glu Pro Ala Glu Arg Pro Lys Leu Thr Leu Lys Asn 10 Val Ile Gly Asp Leu Val Ala Leu Ile Gln Asn Val Leu Phe Asn Pro 25 Asp Phe Ser Val Phe Val Ala Pro Leu Leu Trp Leu Ala Asp Ser Ile 40 Val Ile Lys Val Ile Ile Gly Thr Val Ser Tyr Thr Asp Ile Asp Phe Ser Ser Tyr Met Gln Gln Ile Phe Lys Ile Arg Gln Gly Glu Leu Asp 70 Tyr Ser Asn Ile Phe Gly Asp Thr Gly Pro Leu Val Tyr Pro Ala Gly His Val His Ala Tyr Ser Val Leu Ser Trp Tyr Ser Asp Gly Gly Glu 105 Asp Val Ser Phe Val Gln Gln Ala Phe Gly Trp Leu Tyr Leu Gly Cys 120 Leu Leu Leu Ser Ile Ser Ser Tyr Phe Phe Ser Gly Leu Gly Lys Ile 135 140 Pro Pro Val Tyr Phe Val Leu Leu Val Ala Ser Lys Arg Leu His Ser 150 155 Ile Phe Val Leu Arg Leu Phe Asn Asp Cys Leu Thr Thr Phe Leu Met 165 170

Leu Ala Thr Ile Ile Ile Leu Gln Gln Ala Ser Ser Trp Arg Lys Asp

```
200
                                                 205
Leu Ala Ile Ser Val Lys Met Asn Ala Leu Leu Tyr Leu Pro Ala Phe
                        215
                                             220
Leu Leu Ile Tyr Leu Ile Cys Asp Glu Asn Leu Ile Lys Ala Leu
                    230
                                         235
Ala Pro Val Leu Val Leu Ile Leu Val Gln Val Gly Val Gly Tyr Ser
                245
                                     250
Phe Ile Leu Pro Leu His Tyr Asp Asp Gln Ala Asn Glu Ile Arg Ser
                                265
Ala Tyr Phe Arg Gln Ala Phe Asp Phe Ser Arg Gln Phe Leu Tyr Lys
                            280
Trp Thr Val Asn Trp Arg Phe Leu Ser Gln Glu Thr Phe Asn Asn Val
                        295
His Phe His Gln Leu Leu Phe Ala Leu His Ile Ile Thr Leu Val Leu
                    310
                                         315
Phe Ile Leu Lys Phe Leu Ser Pro Lys Asn Ile Gly Lys Pro Leu Gly
                325
                                    330
Arg Phe Val Leu Asp Ile Phe Lys Phe Trp Lys Pro Thr Leu Ser Pro
            340
                                345
                                                     350
Thr Asn Ile Ile Asn Asp Pro Glu Arg Ser Pro Asp Phe Val Tyr Thr
                            360
                                                 365
Val Met Ala Thr Thr Asn Leu Ile Gly Val Leu Phe Ala Arg Ser Leu
                        375
                                             380
His Tyr Gln Phe Leu Ser Trp Tyr Ala Phe Ser Leu Pro Tyr Leu Leu
                   390
                                        395
Tyr Lys Ala Arg Leu Asn Phe Ile Ala Ser Ile Ile Val Tyr Ala Ala
               405
                                    410
His Glu Tyr Cys Trp Leu Val Phe Pro Ala Thr Glu Gln Ser Ser Ala
            420
                                425
Leu Leu Val Ser Ile Leu Leu Leu Ile Leu Ile Leu Ile Phe Thr Asn
                            440
Glu Gln Leu Phe Pro Ser Gln Ser Val Pro Ala Glu Lys Lys Asn Thr
                        455
<210> 36
<211> 418
<212> PRT
<213> Pichia pastoris
<220>
<221> MUTAGEN
<222> (209)...(223)
<223> Xaa is a variable amino acid
<220>
<221> MOD RES
<222> (235)...(246)
<223> Xaa is a variable amino acid
<400> 36
Arg Pro Lys Leu Thr Leu Lys Asn Val Ile Gly Asp Leu Val Ala Leu
                5
                                    10
Ile Gln Asn Val Leu Phe Asn Pro Asp Phe Ser Val Phe Val Ala Pro
           20
                                25
Leu Leu Trp Leu Ala Asp Ser Ile Val Ile Lys Val Ile Ile Gly Thr
```

180 185 190 Gly Thr Thr Ile Pro Leu Ser Val Pro Asp Ala Ala Asp Thr Tyr Ser

```
40
Val Ser Tyr Thr Asp Ile Asp Phe Ser Ser Tyr Met Gln Gln Ile Phe
                       55
Lys Ile Arg Gln Gly Glu Leu Asp Tyr Ser Asn Ile Phe Gly Asp Thr
                                      75
Gly Pro Leu Val Tyr Pro Ala Gly His Val His Ala Tyr Ser Val Leu
               85
                                   90
Ser Trp Tyr Ser Asp Gly Gly Glu Asp Val Ser Phe Val Gln Gln Ala
                               105
Phe Gly Trp Leu Tyr Leu Gly Cys Leu Leu Ser Ile Ser Ser Tyr
                           120
Phe Phe Ser Gly Leu Gly Lys Ile Pro Pro Val Tyr Phe Val Leu Leu
                       135
                                          140
Val Ala Ser Lys Arg Leu His Ser Ile Phe Val Leu Arg Leu Phe Asn
                   150
Asp Cys Leu Thr Thr Phe Leu Met Leu Ala Thr Ile Ile Ile Leu Gln
                                   170
Gln Ala Ser Ser Trp Arg Lys Asp Gly Thr Thr Ile Pro Leu Ser Val
                              185
                                                  190
Pro Asp Ala Ala Asp Thr Tyr Ser Leu Ala Ile Ser Val Lys Met Asn
                           200
215
Asp Glu Asn Leu Ile Lys Ala Leu Ala Pro Xaa Xaa Xaa Xaa Xaa Xaa
                   230
                                      235
Xaa Xaa Xaa Xaa Xaa Tyr Ser Phe Ile Leu Pro Leu His Tyr Asp
               245
                                                      255
                                  250
Asp Gln Ala Asn Glu Ile Arg Ser Ala Tyr Phe Arg Gln Ala Phe Asp
                              265
Phe Ser Arg Gln Phe Leu Tyr Lys Trp Thr Val Asn Trp Arg Phe Leu
                         280
                                             285
Ser Gln Glu Thr Phe Asn Asn Val His Phe His Gln Leu Leu Phe Ala
                      295
Leu His Ile Ile Thr Leu Val Leu Phe Ile Leu Lys Phe Leu Ser Pro
                   310
                                      315
Lys Asn Ile Gly Lys Pro Leu Gly Arg Phe Val Leu Asp Ile Phe Lys
                                  330
Phe Trp Lys Pro Thr Leu Ser Pro Thr Asn Ile Ile Asn Pro Asp Phe
                              345
Val Tyr Thr Val Met Ala Thr Thr Asn Leu Ile Gly Val Leu Phe Ala
                          360
                                              365
Arg Ser Leu His Tyr Gln Phe Leu Ser Trp Tyr Ala Phe Ser Leu Pro
                       375
                                          380
Tyr Leu Leu Tyr Lys Ala Arg Leu Asn Phe Ile Ala Ser Ile Ile Val
                  390
                                      395
Tyr Ala Ala His Glu Tyr Cys Trp Leu Val Phe Pro Ala Thr Glu Gln
               405
                               410
Ser Ser
```

```
<210> 37
```

<211> 398

<212> PRT

<213> Saccharomyces cerevisiae

<400> 37

Arg Pro Pro Leu Asp Leu Trp Gln Asp Leu Lys Asp Gly Val Arg Tyr

```
Val Ile Phe Asp Cys Arg Ala Asn Leu Ile Val Met Pro Leu Leu Ile
                               25
Leu Phe Glu Ser Met Leu Cys Lys Ile Ile Ile Lys Lys Val Ala Tyr
                           40
Thr Glu Ile Asp Tyr Lys Ala Tyr Met Glu Gln Ile Glu Met Ile Gln
                       5.5
Leu Asp Gly Met Leu Asp Tyr Ser Gln Val Ser Gly Gly Thr Gly Pro
                   70
                                       75
Leu Val Tyr Pro Ala Gly His Val Leu Ile Tyr Lys Met Met Tyr Trp
                                   90
Leu Thr Glu Gly Met Asp His Val Glu Arg Gly Gln Val Phe Phe Arg
                               105
Tyr Leu Tyr Leu Leu Thr Leu Ala Leu Gln Met Ala Cys Tyr Tyr Leu
       115
                           120
Leu His Leu Pro Pro Trp Cys Val Val Leu Ala Cys Leu Ser Lys Arg
                       135
Leu His Ser Ile Tyr Val Leu Arg Leu Phe Asn Asp Cys Phe Thr Thr
                   150
                                       155
Leu Phe Met Val Val Thr Val Leu Gly Ala Ile Val Ala Ser Arg Cys
               165
                                   170
His Gln Arg Pro Lys Leu Lys Ser Leu Ala Leu Val Ile Ser Ala
           180
                               185
Thr Tyr Ser Met Ala Val Ser Ile Lys Met Asn Ala Leu Leu Tyr Phe
                          200
       195
                                               205
Pro Ala Met Met Ile Ser Leu Phe Ile Leu Asn Asp Ala Asn Val Ile
                      215
                                        220
Leu Thr Leu Leu Asp Leu Val Ala Met Ile Ala Trp Gln Val Ala Val
       230
                                       235
Ala Val Pro Phe Leu Arg Ser Phe Pro Gln Gln Tyr Leu His Cys Ala
            245
                                  250
Phe Asn Phe Gly Arg Lys Phe Met Tyr Gln Trp Ser Ile Asn Trp Gln
                              265
Met Met Asp Glu Glu Ala Phe Asn Asp Lys Arg Phe His Leu Ala Leu
                           280
Leu Ile Ser His Leu Ile Ala Leu Thr Thr Leu Phe Val Thr Arg Tyr
                       295
                                           300
Pro Arg Ile Leu Pro Asp Leu Trp Ser Ser Leu Cys His Pro Leu Arg
                  310
                                       315
Lys Asn Ala Val Leu Asn Ala Asn Pro Ala Lys Thr Ile Pro Phe Val
              325
                                   330
Leu Ile Ala Ser Asn Phe Ile Gly Val Leu Phe Ser Arg Ser Leu His
                               345
                                                   350
Tyr Gln Phe Leu Ser Trp Tyr His Trp Thr Leu Pro Ile Leu Ile Phe
       355
                           360
                                               365
Trp Ser Gly Met Pro Phe Phe Val Gly Pro Ile Trp Tyr Val Leu His
                      375
                                           380
Glu Trp Cys Trp Asn Ser Tyr Pro Pro Asn Ser Gln Ala Ser
                   390
                                       395
```

```
<210> 38
```

<211> 387

<212> PRT

<213> Pichia pastoris

<220>

<221> MOD RES

```
<222> (183)...(197)
<223> Xaa is a variable amino acid
<220>
<221> MOD RES
<222> (209)...(220)
<223> Xaa is a variable amino acid
<400> 38
Ser Val Phe Val Ala Pro Leu Leu Trp Leu Ala Asp Ser Ile Val Ile
Lys Val Ile Ile Gly Thr Val Ser Tyr Thr Asp Ile Asp Phe Ser Ser
                              25
Tyr Met Gln Gln Ile Phe Lys Ile Arg Gln Gly Glu Leu Asp Tyr Ser
Asn Ile Phe Gly Asp Thr Gly Pro Leu Val Tyr Pro Ala Gly His Val
                       55
His Ala Tyr Ser Val Leu Ser Trp Tyr Ser Asp Gly Glu Asp Val
                   70
                                     .75
Ser Phe Val Gln Gln Ala Phe Gly Trp Leu Tyr Leu Gly Cys Leu Leu
                                 90
Leu Ser Ile Ser Ser Tyr Phe Phe Ser Gly Leu Gly Lys Ile Pro Pro
           100
                              105
Val Tyr Phe Val Leu Leu Val Ala Ser Lys Arg Leu His Ser Ile Phe
                          120
                                             125
Val Leu Arg Leu Phe Asn Asp Cys Leu Thr Thr Phe Leu Met Leu Ala
                    135
Thr Ile Ile Ile Leu Gln Gln Ala Ser Ser Trp Arg Lys Asp Gly Thr
                  150
                                      155
Thr Ile Pro Leu Ser Val Pro Asp Ala Ala Asp Thr Tyr Ser Leu Ala
                                 170
Ile Ser Val Lys Met Asn Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa
                              185
Xaa Xaa Xaa Xaa Cys Asp Glu Asn Leu Ile Lys Ala Leu Ala Pro
                          200
                                             205
215
                                          220
Leu Pro Leu His Tyr Asp Asp Gln Ala Asn Glu Ile Arg Ser Ala Tyr
                  230
                                      235
Phe Arg Gln Ala Phe Asp Phe Ser Arg Gln Phe Leu Tyr Lys Trp Thr
               245
                                  250
Val Asn Trp Arg Phe Leu Ser Gln Glu Thr Phe Asn Asn Val His Phe
           260
                              265
His Gln Leu Leu Phe Ala Leu His Ile Ile Thr Leu Val Leu Phe Ile
                          280
                                              285
Pro Leu Gly Arg Phe Val Leu Asp Ile Phe Lys Phe Trp Lys Pro Thr
                      295
                                          300
Leu Ser Pro Thr Asn Ile Ile Asn Asp Pro Glu Arg Ser Pro Asp Phe
                   310
                                      315
Val Tyr Thr Val Met Ala Thr Thr Asn Leu Ile Gly Val Leu Phe Ala
               325
                                  330
Arg Ser Leu His Tyr Gln Phe Leu Ser Trp Tyr Ala Phe Ser Leu Pro
                              345
Tyr Leu Leu Tyr Lys Ala Arg Leu Asn Phe Ile Ala Ser Ile Ile Val
                          360
                                              365
Tyr Ala Ala His Glu Tyr Cys Trp Leu Val Phe Pro Ala Thr Glu Gln
                      375
                                          380
Ser Ser Ala
```

385

```
<210> 39
<211> 373
<212> PRT
<213> Neurospora crassa
<400> 39
Ser Lys Leu Ile Pro Pro Ala Leu Phe Leu Val Asp Ala Leu Leu Cys
                                    10
Gly Leu Ile Ile Trp Lys Val Pro Tyr Thr Glu Ile Asp Trp Ala Ala
                                25
Tyr Met Glu Gln Val Ser Gln Ile Leu Ser Gly Glu Arg Asp Tyr Thr
Lys Val Arg Gly Gly Thr Gly Pro Leu Val Tyr Pro Ala Ala His Val
Tyr Ile Tyr Thr Gly Leu Tyr His Leu Thr Asp Glu Gly Arg Asn Ile
                                        75
Leu Leu Ala Gln Gln Leu Phe Ala Gly Leu Tyr Met Val Thr Leu Ala
                                    90
Val Val Met Gly Cys Tyr Trp Gln Ala Lys Ala Pro Pro Tyr Leu Phe
            100
                               105
                                                    110
Pro Leu Leu Thr Leu Ser Lys Arg Leu His Ser Ile Phe Val Leu Arg
                            120
                                                125
Cys Phe Asn Asp Cys Phe Ala Val Leu Phe Leu Trp Leu Ala Ile Phe
                        135
                                            140
Phe Phe Gln Arg Arg Asn Trp Gln Ala Gly Ala Leu Leu Tyr Thr Leu
                   150
                                       155
Gly Leu Gly Val Lys Met Thr Leu Leu Leu Ser Leu Pro Ala Val Gly
               165
                                   170
Ile Val Leu Phe Leu Gly Ser Gly Ser Phe Val Thr Thr Leu Gln Leu
                               185
Val Ala Thr Met Gly Leu Val Gln Ile Leu Ile Gly Val Pro Phe Leu
                            200
                                                205
Ala His Tyr Pro Thr Glu Tyr Leu Ser Arg Ala Phe Glu Leu Ser Arg
                        215
                                            220
Gln Phe Phe Phe Lys Trp Thr Val Asn Trp Arg Phe Val Gly Glu Glu
                   230
                                       235
Ile Phe Leu Ser Lys Gly Phe Ala Leu Thr Leu Leu Ala Leu His Val
               245
                                    250
                                                        255
Leu Val Leu Gly Ile Phe Ile Thr Thr Arg Trp Ile Lys Pro Ala Arg
                                265
Lys Ser Leu Val Gln Leu Ile Ser Pro Val Leu Leu Ala Gly Lys Pro
                            280
                                                285
Pro Leu Thr Val Pro Glu His Arg Ala Ala Ala Arg Asp Val Thr Pro
                        295
                                            300
Arg Tyr Ile Met Thr Thr Ile Leu Ser Ala Asn Ala Val Gly Leu Leu
                    310
                                        315
Phe Ala Arg Ser Leu His Tyr Gln Phe Tyr Ala Tyr Val Ala Trp Ser
                325
                                    330
Thr Pro Phe Leu Leu Trp Arg Ala Gly Leu His Pro Val Leu Val Tyr
                                345
Leu Leu Trp Ala Val His Glu Trp Ala Trp Asn Val Phe Pro Ser Thr
       355
                            360
                                                365
Pro Ala Ser Ser Ala
    370
```

```
<210> 40
<211> 374
<212> PRT
<213> Pichia pastoris
<220>
<221> MOD RES
<222> (160)...(174)
<223> Xaa is a variable amino acid
<220>
<221> MOD RES
<222> (186)...(197)
<223> Xaa is a variable amino acid
<400> 40
Ser Tyr Thr Asp Ile Asp Phe Ser Ser Tyr Met Gln Gln Ile Phe Lys
                                   10
Ile Arg Gln Gly Glu Leu Asp Tyr Ser Asn Ile Phe Gly Asp Thr Gly
                               25
                                                  30
Pro Leu Val Tyr Pro Ala Gly His Val His Ala Tyr Ser Val Leu Ser
                           40
                                              45
Trp Tyr Ser Asp Gly Gly Glu Asp Val Ser Phe Val Gln Gln Ala Phe
Gly Trp Leu Tyr Leu Gly Cys Leu Leu Leu Ser Ile Ser Ser Tyr Phe
                   70
                                      75
Phe Ser Gly Leu Gly Lys Ile Pro Pro Val Tyr Phe Val Leu Leu Val
                                  90
Ala Ser Lys Arg Leu His Ser Ile Phe Val Leu Arg Leu Phe Asn Asp
           100
                              105
Cys Leu Thr Thr Phe Leu Met Leu Ala Thr Ile Ile Leu Gln Gln
                          120
Ala Ser Ser Trp Arg Lys Asp Gly Thr Thr Ile Pro Leu Ser Val Pro
                      135
                                          140
Asp Ala Ala Asp Thr Tyr Ser Leu Ala Ile Ser Val Lys Met Asn Xaa
                  150
                                      155
165
                                  170
Glu Asn Leu Ile Lys Ala Leu Ala Pro Xaa Xaa Xaa Xaa Xaa Xaa
           180
                               185
Xaa Xaa Xaa Xaa Tyr Ser Phe Ile Leu Pro Leu His Tyr Asp Asp
                           200
                                              205
Gln Ala Asn Glu Ile Arg Ser Ala Tyr Phe Arg Gln Ala Phe Asp Phe
                       215
                                          220
Ser Arg Gln Phe Leu Tyr Lys Trp Thr Val Asn Trp Arg Phe Leu Ser
                   230
                                      235
Gln Glu Thr Phe Asn Asn Val His Phe His Gln Leu Leu Phe Ala Leu
               245
                                  250
His Ile Ile Thr Leu Val Leu Phe Ile Leu Lys Phe Leu Ser Pro Lys
           260
                              265
Asn Ile Gly Lys Pro Leu Gly Arg Phe Val Leu Asp Ile Phe Lys Phe
       275
                           280
                                              285
Trp Lys Pro Thr Leu Ser Pro Thr Asn Ile Ile Asn Asp Pro Glu Arg
                      295
                                          300
Ser Pro Asp Phe Val Tyr Thr Val Met Ala Thr Thr Asn Leu Ile Gly
                                      315
                   310
Val Leu Phe Ala Arg Ser Leu His Tyr Gln Phe Leu Ser Trp Tyr Ala
```

```
325
                                    330
Phe Ser Leu Pro Tyr Leu Leu Tyr Lys Ala Arg Leu Asn Phe Ile Ala
                                345
Ser Ile Ile Val Tyr Ala Ala His Glu Tyr Cys Trp Leu Val Phe Pro
                            360
Ala Thr Glu Gln Ser Ser
    370
<210> 41
<211> 355
<212> PRT
<213> Schizosaccharomyces pombe
<400> 41
Leu Leu Leu Glu Ile Pro Phe Val Phe Ala Ile Ile Ser Lys Val
Pro Tyr Thr Glu Ile Asp Trp Ile Ala Tyr Met Glu Gln Val Asn Ser
                                25
Phe Leu Leu Gly Glu Arg Asp Tyr Lys Ser Leu Val Gly Cys Thr Gly
                            40
                                                4.5
Pro Leu Val Tyr Pro Gly Gly His Val Phe Leu Tyr Thr Leu Leu Tyr
Tyr Leu Thr Asp Gly Gly Thr Asn Ile Val Arg Ala Gln Tyr Ile Phe
                    70
                                        75
Ala Phe Val Tyr Trp Ile Thr Thr Ala Ile Val Gly Tyr Leu Phe Lys
                85
                                    90
Ile Val Arg Ala Pro Phe Tyr Ile Tyr Val Leu Leu Ile Leu Ser Lys
                               105
Arg Leu His Ser Ile Phe Ile Leu Arg Leu Phe Asn Asp Gly Phe Asn
        115
                           120
Ser Leu Phe Ser Ser Leu Phe Ile Leu Ser Ser Cys Lys Lys Trp
                       135
Val Arg Ala Ser Ile Leu Leu Ser Val Ala Cys Ser Val Lys Met Ser
                                        155
Ser Leu Leu Tyr Val Pro Ala Tyr Leu Val Leu Leu Gln Ile Leu
                                    170
Gly Pro Lys Lys Thr Trp Met His Ile Phe Val Ile Ile Ile Val Gln
                                185
Ile Leu Phe Ser Ile Pro Phe Leu Ala Tyr Phe Trp Ser Tyr Trp Thr
                            200
Gln Ala Phe Asp Phe Gly Arg Ala Phe Asp Tyr Lys Trp Thr Val Asn
                       215
                                            220
Trp Arg Phe Ile Pro Arg Ser Ile Phe Glu Ser Thr Ser Phe Ser Thr
                   230
                                        235
Ser Ile Leu Phe Leu His Val Ala Leu Leu Val Ala Phe Thr Cys Lys
               245
                                    250
His Trp Asn Lys Leu Ser Arg Ala Thr Pro Phe Ala Met Val Asn Ser
           260
                                265
Met Leu Thr Leu Lys Pro Leu Pro Lys Leu Gln Leu Ala Thr Pro Asn
                            280
Phe Ile Phe Thr Ala Leu Ala Thr Ser Asn Leu Ile Gly Ile Leu Cys
                       295
Ala Arg Ser Leu His Tyr Gln Phe Tyr Ala Trp Phe Ala Trp Tyr Ser
                   310
                                        315
Pro Tyr Leu Cys Tyr Gln Ala Ser Phe Pro Ala Pro Ile Val Ile Gly
                                    330
Leu Trp Met Leu Gln Glu Tyr Ala Trp Asn Val Phe Pro Ser Thr Lys
```

```
340
                            345
                                                  350
Leu Ser Ser
       355
<210> 42
<211> 390
<212> PRT
<213> Pichia pastoris
<220>
<221> MOD RES
<222> (176)...(190)
<223> Xaa is a variable amino acid
<220>
<221> MOD RES
<222> (202)...(213)
<223> Xaa is a variable amino acid
<400> 42
Leu Trp Leu Ala Asp Ser Ile Val Ile Lys Val Ile Ile Gly Thr Val
               5
                               10
Ser Tyr Thr Asp Ile Asp Phe Ser Ser Tyr Met Gln Gln Ile Phe Lys
                              25
Ile Arg Gln Gly Glu Leu Asp Tyr Ser Asn Ile Phe Gly Asp Thr Gly
                          40
Pro Leu Val Tyr Pro Ala Gly His Val His Ala Tyr Ser Val Leu Ser
                       55
Trp Tyr Ser Asp Gly Glu Asp Val Ser Phe Val Gln Gln Ala Phe
                  70
Gly Trp Leu Tyr Leu Gly Cys Leu Leu Leu Ser Ile Ser Ser Tyr Phe
Phe Ser Gly Leu Gly Lys Ile Pro Pro Val Tyr Phe Val Leu Leu Val
                              105
Ala Ser Lys Arg Leu His Ser Ile Phe Val Leu Arg Leu Phe Asn Asp
                          120
Cys Leu Thr Thr Phe Leu Met Leu Ala Thr Ile Ile Ile Leu Gln Gln
                      135
                                         140
Ala Ser Ser Trp Arg Lys Asp Gly Thr Thr Ile Pro Leu Ser Val Pro
                  150
                                      155
Asp Ala Ala Asp Thr Tyr Ser Leu Ala Ile Ser Val Lys Met Asn Xaa
               165
                                 170
180
                              185
                                                 190
Glu Asn Leu Ile Lys Ala Leu Ala Pro Xaa Xaa Xaa Xaa Xaa Xaa
                          200
                                              205
Xaa Xaa Xaa Xaa Tyr Ser Phe Ile Leu Pro Leu His Tyr Asp Asp
                      215
                                          220
Gln Ala Asn Glu Ile Arg Ser Ala Tyr Phe Arg Gln Ala Phe Asp Phe
                  230
                                      235
Ser Arg Gln Phe Leu Tyr Lys Trp Thr Val Asn Trp Arg Phe Leu Ser
               245
                                  250
Gln Glu Thr Phe Asn Asn Val His Phe His Gln Leu Leu Phe Ala Leu
                              265
                                                 270
His Ile Ile Thr Leu Val Leu Phe Ile Leu Lys Phe Leu Ser Pro Lys
                          280
                                             285
Asn Ile Gly Lys Pro Leu Gly Arg Phe Val Leu Asp Ile Phe Lys Phe
```

```
295
Trp Lys Pro Thr Leu Ser Pro Thr Asn Ile Ile Asn Asp Pro Glu Arg
                    310
                                        315
Ser Pro Asp Phe Val Tyr Thr Val Met Ala Thr Thr Asn Leu Ile Gly
                325
                                    330
Val Leu Phe Ala Arg Ser Leu His Tyr Gln Phe Leu Ser Trp Tyr Ala
                                345
Phe Ser Leu Pro Tyr Leu Leu Tyr Lys Ala Arg Leu Asn Phe Ile Ala
                            360
                                                365
Ser Ile Ile Val Tyr Ala Ala His Glu Tyr Cys Trp Leu Val Phe Pro
                        375
                                            380
Ala Thr Glu Gln Ser Ser
<210> 43
<211> 363
<212> PRT
<213> Arabidopsis thaliana
<400> 43
Leu Ile Leu Ala Asp Ala Ile Leu Val Ala Leu Ile Ile Ala Tyr Val
    . 5
                                   10
Pro Tyr Thr Lys Ile Asp Trp Asp Ala Tyr Met Ser Gln Val Ser Gly
                                25
Phe Leu Gly Gly Glu Arg Asp Tyr Gly Asn Leu Lys Gly Asp Thr Gly
                           40
                                                45
Pro Leu Val Tyr Pro Ala Gly Phe Leu Tyr Val Tyr Ser Ala Val Gln
                        55
                                            60
Asn Leu Thr Gly Gly Glu Val Tyr Pro Ala Gln Ile Leu Phe Gly Val
                   70
                                        75
Leu Tyr Ile Val Asn Leu Gly Ile Val Leu Ile Ile Tyr Val Lys Thr
                                    90
Asp Val Val Pro Trp Trp Ala Leu Ser Leu Leu Cys Leu Ser Lys Arg
                                105
Ile His Ser Ile Phe Val Leu Arg Leu Phe Asn Asp Cys Phe Ala Met
                            120
Thr Leu Leu His Ala Ser Met Ala Leu Phe Leu Tyr Arg Lys Trp His
                        135
                                            140
Leu Gly Met Leu Val Phe Ser Gly Ala Val Ser Val Lys Met Asn Val
                   150
                                        155
Leu Leu Tyr Ala Pro Thr Leu Leu Leu Leu Leu Leu Lys Ala Met Asn
               165
                                    170
Ile Ile Gly Val Val Ser Ala Leu Ala Gly Ala Ala Leu Ala Gln Ile
           180
                                185
Leu Val Gly Leu Pro Phe Leu Ile Thr Tyr Pro Val Ser Tyr Ile Ala
        195
                            200
                                                205
Asn Ala Phe Asp Leu Gly Arg Val Phe Ile His Phe Trp Ser Val Asn
                        215
                                            220
Phe Lys Phe Val Pro Glu Arg Val Phe Val Ser Lys Glu Phe Ala Val
                   230
                                        235
Cys Leu Leu Ile Ala His Leu Phe Leu Leu Val Ala Phe Ala Asn Tyr
               245
                                    250
Lys Trp Cys Lys His Glu Gly Gly Ile Ile Gly Phe Met Arg Ser Arg
                               265
                                                    270
His Phe Phe Leu Thr Leu Pro Ser Ser Leu Ser Phe Ser Asp Val Ser
                           280
                                                285
Ala Ser Arg Ile Ile Thr Lys Glu His Val Val Thr Ala Met Phe Val
```

```
295
Gly Asn Phe Ile Gly Ile Val Phe Ala Arg Ser Leu His Tyr Gln Phe
                    310
                                         315
Tyr Ser Trp Tyr Phe Tyr Ser Leu Pro Tyr Leu Leu Trp Arg Thr Pro
                325
                                     330
Phe Pro Thr Trp Leu Arg Leu Ile Met Phe Leu Gly Ile Glu Leu Cys
                                 345
Trp Asn Val Tyr Pro Ser Thr Pro Ser Ser Ser
                             360
<210> 44
<211> 428
<212> DNA
<213> Kluyveromyces lactis
<400> 44
tttgtttaca agctgatacc aacgaacatg aatacaccgg caggtttact gaagattggc 60
aaagctaacc ttttacatcc ttttaccgat gctgtattca gtgcgatgag agtaaacgca 120
gaacaaattg catacatttt acttgttacc aattacattg gagtactatt tgctcgatca 180
ttacactacc aattcctatc ttggtaccat tggacgttac cagtactatt gaattgggcc 240
aatgttccgt atccgctatg tgtgctatgg tacctaacac atgagtggtg ctggaacagc 300
tatccgccaa acgctactgc atccacactg ctacacgcgt gtaacacata ctgttattgg 360
ctgtattctt aagaggaccc gcaaactcga aaagtggtga taacgaaaca acacacgaga 420
aagctgag
<210> 45
<211> 141
<212> PRT
<213> Kluyveromyces lactis
<400> 45
Phe Val Tyr Lys Leu Ile Pro Thr Asn Met Asn Thr Pro Ala Gly Leu
                                     10
Leu Lys Ile Gly Lys Ala Asn Leu Leu His Pro Phe Thr Asp Ala Val
                                25
Phe Ser Ala Met Arg Val Asn Ala Glu Gln Ile Ala Tyr Ile Leu Leu
Val Thr Asn Tyr Ile Gly Val Leu Phe Ala Arg Ser Leu His Tyr Gln
                        5.5
Phe Leu Ser Trp Tyr His Trp Thr Leu Pro Val Leu Leu Asn Trp Ala
                    70
                                         7.5
Asn Val Pro Tyr Pro Leu Cys Val Leu Trp Tyr Leu Thr His Glu Trp
                85
                                     90
Cys Trp Asn Ser Tyr Pro Pro Asn Ala Thr Ala Ser Thr Leu Leu His
                                105
Ala Cys Asn Thr Tyr Cys Tyr Trp Leu Tyr Ser Glu Asp Pro Gln Thr
                            120
Arg Lys Val Val Ile Thr Lys Gln His Thr Arg Lys Leu
                        135
<210> 46
<211> 118
<212> PRT
<213> Kluyveromyces lactis
<400> 46
```

<210> 47

<211> 117

<212> PRT

<213> Saccharomyces cerevisiae

<400> 47

 Ser
 Leu
 Cys
 His
 Pro
 Leu
 Arg
 Lys
 Asn
 Ala
 Val
 Leu
 Asn
 Ala
 Ala</th

<210> 48

<211> 113

<212> PRT

<213> Kluyveromyces lactis

<400> 48

 Phe
 Thr
 Asp
 Ala
 Val
 Phe
 Ser
 Ala
 Met
 Arg
 Val
 Asn
 Ala
 Glu
 Gln
 Ile

 Ala
 Tyr
 Ile
 Leu
 Leu
 Leu
 Val
 Tyr
 Ile
 Gly
 Val
 Leu
 Phe
 Ala
 Arg

 Ser
 Leu
 His
 Tyr
 Gln
 Phe
 Leu
 Ser
 Tyr
 Tyr
 His
 Tyr
 Thr
 Leu
 Pro
 Val

 Leu
 Leu
 Asn
 Tyr
 Pro
 Tyr
 Pro
 Leu
 Cys
 Val
 Leu
 Tyr
 Tyr
 Tyr
 Pro
 Pro
 Asn
 Ala
 Thr
 Ala

 Leu
 Tyr
 His
 Glu
 Tyr
 Cys
 Tyr
 Pro
 Pro
 Pro
 Asn
 Ala
 Thr
 Ala

 Leu
 Tyr
 His
 Glu
 Tyr
 Pyr
 P

```
Ser Thr Leu Leu His Ala Cys Asn Thr Tyr Cys Tyr Trp Leu Tyr Ser
Glu Asp Pro Gln Thr Arg Lys Val Val Ile Thr Lys Gln His Thr Arg
                                105
Lys
<210> 49
<211> 106
<212> PRT
<213> Arabidopsis thaliana
<400> 49
Phe Ser Asp Val Ser Ala Ser Arg Ile Ile Thr Lys Glu His Val Val
 1
Thr Ala Met Phe Val Gly Asn Phe Ile Gly Ile Val Phe Ala Arg Ser
                                25
Leu His Tyr Gln Phe Tyr Ser Trp Tyr Phe Tyr Ser Leu Pro Tyr Leu
                            40
Leu Trp Arg Thr Pro Phe Pro Thr Trp Leu Arg Leu Ile Met Phe Leu
                        55
                                            60
Gly Ile Glu Leu Cys Trp Asn Val Tyr Pro Ser Thr Pro Ser Ser Ser
                    70
                                        75
Gly Leu Leu Cys Leu His Leu Ile Ile Leu Val Gly Leu Trp Leu
                85
                                    90
Ala Pro Ser Val Asp Pro Tyr Gln Leu Lys
            100
                                105
<210> 50
<211> 1668
<212> DNA
<213> Saccharomyces cerevisiae
atgaattgca aggcggtaac cattagttta ttactgttgt tatttttaac aagagtatat 60
attcagccga cattctcgtt aatttcagat tgcgatgaaa cttttaatta ttgggaacca 120
ttaaatttat tggtacgtgg atttggtaaa caaacctggg aatattcacc cgagtattct 180
attagateat gggetttett attacetttt taetgtatte tttateeagt aaacaaattt 240
actgacctag aaagtcattg gaactttttc atcacaagag catgcttagg cttttttagt 300
tttatcatgg aatttaaact acatcgtgaa attgcaggca gcttggcatt gcaaatcgca 360
aatatttgga ttattttcca attgtttaat ccgggctggt tccatgcatc tgtggaatta 420
ttgccttctg ccgttgccat gttgttgtat gtaggtgcca ccagacactc tctacgctat 480
ctgtccactg ggtctacttc taactttacg aaaagtttag cgtacaattt cctggctagt 540
atactagget ggecatttgt tttaatttta agettgecat tatgtttaca ttacetttte 600
aaccatagaa ttatttctac catcagaacc gcattcgact gctgtttgat attttcattg 660
actgcatttg ctgtgattgt cactgacagt atattttacg ggaagettge teetgtatea 720
tggaacatct tattttacaa tgtcattaat gcaagtgagg aatctggccc aaatattttc 780
ggggttgage catggtacta ctatccacta aatttgttac tgaatttece actgeetgtg 840
ctagttttag ctattttggg aattttccat ttgagattat ggccattatg ggcatcatta 900
ttcacatgga ttgccgtttt cactcaacaa cctcacaaag aggaaagatt tctctatcca 960
atttacgggt taataacttt gagtgcaagt atcgcctttt acaaagtgtt gaatctattc 1020
aatagaaage egattettaa aaaaggtata aagttgteag ttttattaat tgttgeagge 1080
caggcaatgt cacggatagt ggctttggtg aacaattaca cagctcctat agccgtctac 1140
gagcaatttt cttcactaaa tcaaggtggt gtgaaggcac cggtagtgaa tgtatgtacg 1200
ggacgtgaat ggtatcactt cccaagttct ttcctgctgc cagataatca taggctaaaa 1260
```

tttgttaaat ctggatttga tggtcttctt ccaggtgatt ttccagagag tggttctatt 1320

```
aaagagtggc cgatcactag atgtgattat tttattgaca tcgtcgcccc aataaattta 1440
acaaaagacg ttttcaaccc tctacatctg atggataact ggaataagct ggcatgtgct 1500
gcattcatcg acggtgaaaa ttctaagatt ttgggtagag cattttacgt accggagcca 1560
atcaaccgaa tcatgcaaat agttttacca aaacaatgga atcaagtgta cggtgttcgt 1620
tacattgatt actgtttgtt tgaaaaacca actgagacta ctaattga
<210> 51
<211> 555
<212> PRT
<213> Saccharomyces cerevisiae
<400> 51
Met Asn Cys Lys Ala Val Thr Ile Ser Leu Leu Leu Leu Phe Leu
 1
                                     10
Thr Arg Val Tyr Ile Gln Pro Thr Phe Ser Leu Ile Ser Asp Cys Asp
                                25
Glu Thr Phe Asn Tyr Trp Glu Pro Leu Asn Leu Leu Val Arg Gly Phe
                            40
Gly Lys Gln Thr Trp Glu Tyr Ser Pro Glu Tyr Ser Ile Arg Ser Trp
                        55
                                             60
Ala Phe Leu Pro Phe Tyr Cys Ile Leu Tyr Pro Val Asn Lys Phe
                    70
                                        75
Thr Asp Leu Glu Ser His Trp Asn Phe Phe Ile Thr Arg Ala Cys Leu
                8.5
                                    90
                                                         95
Gly Phe Phe Ser Phe Ile Met Glu Phe Lys Leu His Arg Glu Ile Ala
            100
                                105
                                                     110
Gly Ser Leu Ala Leu Gln Ile Ala Asn Ile Trp Ile Ile Phe Gln Leu
        115
                            120
                                                 125
Phe Asn Pro Gly Trp Phe His Ala Ser Val Glu Leu Leu Pro Ser Ala
                        135
Val Ala Met Leu Leu Tyr Val Gly Ala Thr Arg His Ser Leu Arg Tyr
                    150
                                        155
Leu Ser Thr Gly Ser Thr Ser Asn Phe Thr Lys Ser Leu Ala Tyr Asn
                165
                                    170
Phe Leu Ala Ser Ile Leu Gly Trp Pro Phe Val Leu Ile Leu Ser Leu
                                185
                                                     190
Pro Leu Cys Leu His Tyr Leu Phe Asn His Arg Ile Ile Ser Thr Ile
                            200
                                                 205
Arg Thr Ala Phe Asp Cys Cys Leu Ile Phe Ser Leu Thr Ala Phe Ala
                        215
                                            220
Val Ile Val Thr Asp Ser Ile Phe Tyr Gly Lys Leu Ala Pro Val Ser
                    230
                                        235
Trp Asn Ile Leu Phe Tyr Asn Val Ile Asn Ala Ser Glu Glu Ser Gly
                245
                                    250
Pro Asn Ile Phe Gly Val Glu Pro Trp Tyr Tyr Tyr Pro Leu Asn Leu
            260
                                265
Leu Leu Asn Phe Pro Leu Pro Val Leu Val Leu Ala Ile Leu Gly Ile
       275
                            280
Phe His Leu Arg Leu Trp Pro Leu Trp Ala Ser Leu Phe Thr Trp Ile
                        295
                                            300
Ala Val Phe Thr Gln Gln Pro His Lys Glu Glu Arg Phe Leu Tyr Pro
                    310
                                        315
Ile Tyr Gly Leu Ile Thr Leu Ser Ala Ser Ile Ala Phe Tyr Lys Val
                                    330
                325
                                                         335
Leu Asn Leu Phe Asn Arg Lys Pro Ile Leu Lys Lys Gly Ile Lys Leu
            340
                                345
                                                     350
Ser Val Leu Leu Ile Val Ala Gly Gln Ala Met Ser Arg Ile Val Ala
```

ttcaaaaaga ttagaacttt acctaaggga atgaataaca agaatatata tgataccggt 1380

```
360
Leu Val Asn Asn Tyr Thr Ala Pro Ile Ala Val Tyr Glu Gln Phe Ser
                         375
                                             380
Ser Leu Asn Gln Gly Gly Val Lys Ala Pro Val Val Asn Val Cys Thr
                     390
                                         395
                                                              400
Gly Arg Glu Trp Tyr His Phe Pro Ser Ser Phe Leu Leu Pro Asp Asn
                 405
                                     410
                                                          415
His Arg Leu Lys Phe Val Lys Ser Gly Phe Asp Gly Leu Leu Pro Gly
            420
                                 425
Asp Phe Pro Glu Ser Gly Ser Ile Phe Lys Lys Ile Arg Thr Leu Pro
                             440
Lys Gly Met Asn Asn Lys Asn Ile Tyr Asp Thr Gly Lys Glu Trp Pro
    450
                         455
Ile Thr Arg Cys Asp Tyr Phe Ile Asp Ile Val Ala Pro Ile Asn Leu
                     470
                                         475
Thr Lys Asp Val Phe Asn Pro Leu His Leu Met Asp Asn Trp Asn Lys
                                     490
Leu Ala Cys Ala Ala Phe Ile Asp Gly Glu Asn Ser Lys Ile Leu Gly
            500
                                 505
                                                     510
Arg Ala Phe Tyr Val Pro Glu Pro Ile Asn Arg Ile Met Gln Ile Val
        515
                             520
                                                 525
Leu Pro Lys Gln Trp Asn Gln Val Tyr Gly Val Arg Tyr Ile Asp Tyr
                        535
Cys Leu Phe Glu Lys Pro Thr Glu Thr Thr Asn
545
                    550
<210> 52
<211> 600
<212> DNA
<213> Pichia pastoris
<400> 52
tggccttcct gtctgctcga tacttccttt tacagtaacc aacatacatg ttctccaaca 60
tgctcttgta tgtattggcc tattctatct tgagacttga tatcaacctt ctatggtatt 120
atttcagact gtgatgaagt gttcaactac tgggagccac tcaacttcat gcttagaggg 180
tttggaaaac agacttggga gtattctcca gagtatgcca tccgatcttg gtcctatcta 240
gtgccacttt ggatagcagg ctatccacca ttgttcctgg atatcccttc ttactacttt 300
ttctactttt tcagactact gctggttatt ttttcattgg ttgcagaagt caagttgtac 360
catagtttga agaaaaatgt cagcagtaag atcagtttct ggtaccttct atttacaacc 420
gttgctccag gaatgtctca tagcacgata gccttattac catcctcttt tgctatggtt 480
tgtcacactt ttgccattag atacgtcatt gattacctac aattaccaac attaatgcgc 540
acaatcagag agactgctgc catctcacca gctcacaaac aacaactagc caactctctc 600
<210> 53
<211> 199
<212> PRT
<213> Pichia pastoris
<400> 53
Trp Pro Ser Cys Leu Leu Asp Thr Ser Phe Tyr Ser Asn Gln His Thr
                                    10
Cys Ser Pro Thr Cys Ser Cys Met Tyr Trp Pro Ile Leu Ser Asp Leu
            20
                                25
Ile Ser Thr Phe Tyr Gly Ile Ile Ser Asp Cys Asp Glu Val Phe Asn
                            40
Tyr Trp Glu Pro Leu Asn Phe Met Leu Arg Gly Phe Gly Lys Gln Thr
```

```
55
Trp Glu Tyr Ser Pro Glu Tyr Ala Ile Arg Ser Trp Ser Tyr Leu Val
                    70
Pro Leu Trp Ile Ala Gly Tyr Pro Pro Leu Phe Leu Asp Ile Pro Ser
                                     90
Tyr Tyr Phe Phe Tyr Phe Phe Arg Leu Leu Val Ile Phe Ser Leu
                                105
Val Ala Glu Val Lys Leu Tyr His Ser Leu Lys Lys Asn Val Ser Ser
                            120
Lys Ile Ser Phe Trp Tyr Leu Leu Phe Thr Thr Val Ala Pro Gly Met
                        135
                                             140
Ser His Ser Thr Ile Ala Leu Leu Pro Ser Ser Phe Ala Met Val Cys
                    150
                                        155
His Thr Phe Ala Ile Arg Tyr Val Ile Asp Tyr Leu Gln Leu Pro Thr
                                    170
Leu Met Arg Thr Ile Arg Glu Thr Ala Ala Ile Ser Pro Ala His Lys
            180
                                185
Gln Gln Leu Ala Asn Ser Leu
        195
<210> 54
<211> 140
<212> PRT
<213> Pichia pastoris
<220>
<221> MOD RES
<222> (65)...(71)
<223> Xaa is a variable amino acid
<400> 54
Ile Ser Thr Phe Tyr Gly Ile Ile Ser Asp Cys Asp Glu Val Phe Asn
Tyr Trp Glu Pro Leu Asn Phe Met Leu Arg Gly Phe Gly Lys Gln Thr
Trp Glu Tyr Ser Pro Glu Tyr Ala Ile Arg Ser Trp Ser Tyr Leu Val
                            40
Pro Leu Trp Ile Ala Gly Tyr Pro Pro Leu Phe Leu Asp Ile Pro Ser
Xaa Xaa Xaa Xaa Xaa Xaa Arg Leu Leu Val Ile Phe Ser Leu
                    70
Val Ala Glu Val Lys Leu Tyr His Ser Leu Lys Lys Asn Val Ser Ser
                                    90
Lys Ile Ser Phe Trp Tyr Leu Leu Phe Thr Thr Val Ala Pro Gly Met
           100
                                105
Ser His Ser Thr Ile Ala Leu Leu Pro Ser Ser Phe Ala Met Val Cys
                           120
His Thr Phe Ala Ile Arg Tyr Val Ile Asp Tyr Leu
                        135
<210> 55
<211> 141
<212> PRT
<213> Saccharomyces cerevisiae
<400> 55
```

```
Ile Gln Pro Thr Phe Ser Leu Ile Ser Asp Cys Asp Glu Thr Phe Asn
                                    10
Tyr Trp Glu Pro Leu Asn Leu Leu Val Arg Gly Phe Gly Lys Gln Thr
Trp Glu Tyr Ser Pro Glu Tyr Ser Ile Arg Ser Trp Ala Phe Leu Leu
                            40
Pro Phe Tyr Cys Ile Leu Tyr Pro Val Asn Lys Phe Thr Asp Leu Glu
                        55
Ser His Trp Asn Phe Phe Ile Thr Arg Ala Cys Leu Gly Phe Phe Ser
                    70
Phe Ile Met Glu Phe Lys Leu His Arg Glu Ile Ala Gly Ser Leu Ala
Leu Gln Ile Ala Asn Ile Trp Ile Ile Phe Gln Leu Phe Asn Pro Gly
                                105
Trp Phe His Ala Ser Val Glu Leu Leu Pro Ser Ala Val Ala Met Leu
                            120
Leu Tyr Val Gly Ala Thr Arg His Ser Leu Arg Tyr Leu
                        135
<210> 56
<211> 127
<212> PRT
<213> Pichia pastoris
<220>
<221> MOD RES
<222> (66)...(72)
<223> Xaa is a variable amino acid
<400> 56
Leu Ile Ser Thr Phe Tyr Gly Ile Ile Ser Asp Cys Asp Glu Val Phe
Asn Tyr Trp Glu Pro Leu Asn Phe Met Leu Arg Gly Phe Gly Lys Gln
                                25
Thr Trp Glu Tyr Ser Pro Glu Tyr Ala Ile Arg Ser Trp Ser Tyr Leu
                            40
Val Pro Leu Trp Ile Ala Gly Tyr Pro Pro Leu Phe Leu Asp Ile Pro
Ser Xaa Xaa Xaa Xaa Xaa Xaa Arg Leu Leu Leu Val Ile Phe Ser
                   70
Leu Val Ala Glu Val Lys Leu Tyr His Ser Leu Lys Lys Asn Val Ser
                                    90
               85
Ser Lys Ile Ser Phe Trp Tyr Leu Leu Phe Thr Thr Val Ala Pro Gly
           100
                               105
Met Ser His Ser Thr Ile Ala Leu Leu Pro Ser Ser Phe Ala Met
                           120
<210> 57
<211> 127
<212> PRT
<213> Anopheles gambiae
<400> 57
```

- 31 -

Leu Gln Ser Ala Leu Tyr Ser Ile Ile Ser Asp Cys Asp Glu Thr Tyr

Asn Tyr Trp Glu Pro Leu His Tyr Leu Leu Lys Gly Lys Gly Phe Gln

```
20
Thr Trp Glu Tyr Ser Pro Glu Phe Ala Leu Arg Ser Tyr Ser Tyr Leu
Trp Leu His Gly Leu Pro Ala Lys Val Leu Gln Leu Met Thr Asp Asn
                        55
Gly Val Leu Ile Phe Tyr Phe Val Arg Cys Leu Leu Ala Val Thr Cys
                    70
Ala Leu Leu Glu Tyr Arg Leu Tyr Arg Ile Leu Gly Arg Lys Cys Gly
                8.5
                                    90
Gly Gly Val Ala Ser Leu Trp Leu Leu Phe Gln Leu Thr Ser Ala Gly
                                105
Met Phe Ile Ser Ser Ala Ala Leu Leu Pro Ser Ser Phe Ser Met
                            120
<210> 58
<211> 157
<212> PRT
<213> Pichia pastoris
<220>
<221> MOD RES
<222> (66)...(72)
<223> Xaa is a variable amino acid
<400> 58
Leu Ile Ser Thr Phe Tyr Gly Ile Ile Ser Asp Cys Asp Glu Val Phe
                                    10
Asn Tyr Trp Glu Pro Leu Asn Phe Met Leu Arg Gly Phe Gly Lys Gln
           20
                               25
Thr Trp Glu Tyr Ser Pro Glu Tyr Ala Ile Arg Ser Trp Ser Tyr Leu
                            40
Val Pro Leu Trp Ile Ala Gly Tyr Pro Pro Leu Phe Leu Asp Ile Pro
                        55
Ser Xaa Xaa Xaa Xaa Xaa Xaa Arg Leu Leu Val Ile Phe Ser
                    70
Leu Val Ala Glu Val Lys Leu Tyr His Ser Leu Lys Lys Asn Val Ser
                                    90
Ser Lys Ile Ser Phe Trp Tyr Leu Leu Phe Thr Thr Val Ala Pro Gly
                                105
Met Ser His Ser Thr Ile Ala Leu Leu Pro Ser Ser Phe Ala Met Val
                            120
                                                125
Cys His Thr Phe Ala Ile Arg Tyr Val Ile Asp Tyr Leu Gln Leu Pro
                       135
Thr Leu Met Arg Thr Ile Arg Glu Thr Ala Ala Ile Ser
                    150
                                        155
<210> 59
<211> 154
<212> PRT
<213> Schizosaccharomyces pombe
<400> 59
Leu Thr Ser Ala Ser Phe Arg Val Ile Asp Asp Cys Asp Glu Val Tyr
Asn Tyr Trp Glu Pro Leu His Tyr Leu Leu Tyr Gly Tyr Gly Leu Gln
                                25
```

```
Thr Trp Glu Tyr Ser Pro Glu Tyr Ala Ile Arg Ser Trp Phe Tyr Ile
                            40
Ala Leu His Ala Val Pro Gly Phe Leu Ala Arg Gly Leu Gly Leu Ser
                        55
Arg Leu His Val Phe Tyr Phe Ile Arg Gly Val Leu Ala Cys Phe Ser
                    70
Ala Phe Cys Glu Thr Asn Leu Ile Leu Ala Val Ala Arg Asn Phe Asn
                85
                                    90
Arg Ala Val Ala Leu His Leu Thr Ser Val Leu Phe Val Asn Ser Gly
            100
                                105
Met Trp Ser Ala Ser Thr Ser Phe Leu Pro Ser Ser Phe Ala Met Asn
        115
                            120
Met Val Thr Leu Ala Leu Ser Ala Gln Leu Ser Pro Pro Ser Thr Lys
                        135
Arg Thr Val Lys Val Val Ser Phe Ile Thr
                    150
<210> 60
<211> 141
<212> PRT
<213> Pichia pastoris
<220>
<221> MOD RES
<222> (80)...(86)
<223> Xaa is a variable amino acid
<400> 60
Ser Pro Thr Cys Ser Cys Met Tyr Trp Pro Ile Leu Ser Asp Leu Ile
Ser Thr Phe Tyr Gly Ile Ile Ser Asp Cys Asp Glu Val Phe Asn Tyr
Trp Glu Pro Leu Asn Phe Met Leu Arg Gly Phe Gly Lys Gln Thr Trp
                            40
Glu Tyr Ser Pro Glu Tyr Ala Ile Arg Ser Trp Ser Tyr Leu Val Pro
Leu Trp Ile Ala Gly Tyr Pro Pro Leu Phe Leu Asp Ile Pro Ser Xaa
                    70
Xaa Xaa Xaa Xaa Xaa Arg Leu Leu Leu Val Ile Phe Ser Leu Val
               85
                                    90
Ala Glu Val Lys Leu Tyr His Ser Leu Lys Lys Asn Val Ser Ser Lys
           100
                                105
Ile Ser Phe Trp Tyr Leu Leu Phe Thr Thr Val Ala Pro Gly Met Ser
                           120
His Ser Thr Ile Ala Leu Leu Pro Ser Ser Phe Ala Met
                       135
<210> 61
<211> 143
<212> PRT
<213> Mus musculus
<400> 61
Ala Pro Glu Gly Ser Thr Ala Phe Lys Cys Leu Leu Ser Ala Arg Leu
                                    10
```

Cys Ala Ala Leu Leu Ser Asn Ile Ser Asp Cys Asp Glu Thr Phe Asn

```
Tyr Trp Glu Pro Thr His Tyr Leu Ile Tyr Gly Lys Gly Phe Gln Thr
Trp Glu Tyr Ser Pro Val Tyr Ala Ile Arg Ser Tyr Ala Tyr Leu Leu
                        55
Leu His Ala Trp Pro Ala Ala Phe His Ala Arg Ile Leu Gln Thr Asn
                                        75
                    70
Lys Ile Leu Val Phe Tyr Phe Leu Arg Cys Leu Leu Ala Phe Val Ser
                                    90
Cys Val Cys Glu Leu Tyr Phe Tyr Lys Ala Val Cys Lys Lys Phe Gly
                                105
Leu His Val Ser Arg Met Met Leu Ala Phe Leu Val Leu Ser Thr Gly
                            120
Met Phe Cys Ser Ser Ser Ala Phe Leu Pro Ser Ser Phe Cys Met
                        135
<210> 62
<211> 141
<212> PRT
<213> Pichia pastoris
<220>
<221> MOD RES
<222> (80)...(86)
<223> Xaa is a variable amino acid
<400> 62
Ser Pro Thr Cys Ser Cys Met Tyr Trp Pro Ile Leu Ser Asp Leu Ile
Ser Thr Phe Tyr Gly Ile Ile Ser Asp Cys Asp Glu Val Phe Asn Tyr
           20
Trp Glu Pro Leu Asn Phe Met Leu Arg Gly Phe Gly Lys Gln Thr Trp
                            40
Glu Tyr Ser Pro Glu Tyr Ala Ile Arg Ser Trp Ser Tyr Leu Val Pro
                        55
Leu Trp Ile Ala Gly Tyr Pro Pro Leu Phe Leu Asp Ile Pro Ser Xaa
                                        75
Xaa Xaa Xaa Xaa Xaa Arg Leu Leu Val Ile Phe Ser Leu Val
Ala Glu Val Lys Leu Tyr His Ser Leu Lys Lys Asn Val Ser Ser Lys
                                105
                                                    110
Ile Ser Phe Trp Tyr Leu Leu Phe Thr Thr Val Ala Pro Gly Met Ser
                           120
His Ser Thr Ile Ala Leu Leu Pro Ser Ser Phe Ala Met
                        135
<210> 63
<211> 143
<212> PRT
<213> Homo sapiens
<400> 63
Ala Pro Glu Gly Ser Thr Ala Phe Lys Cys Leu Leu Ser Ala Arg Leu
                5
                                    10
Cys Ala Ala Leu Leu Ser Asn Ile Ser Asp Cys Asp Glu Thr Phe Asn
            20
                                25
```

```
Tyr Trp Glu Pro Thr His Tyr Leu Ile Tyr Gly Glu Gly Phe Gln Thr
Trp Glu Tyr Ser Pro Ala Tyr Ala Ile Arg Ser Tyr Ala Tyr Leu Leu
                        55
Leu His Ala Trp Pro Ala Ala Phe His Ala Arg Ile Leu Gln Thr Asn
                                        75
Lys Ile Leu Val Phe Tyr Phe Leu Arg Cys Leu Leu Ala Phe Val Ser
                                    90
Cys Ile Cys Glu Leu Tyr Phe Tyr Lys Ala Val Cys Lys Lys Phe Gly
            100
                                105
Leu His Val Ser Arg Met Met Leu Ala Phe Leu Val Leu Ser Thr Gly
                            120
Met Phe Cys Ser Ser Ser Ala Phe Leu Pro Ser Ser Phe Cys Met
    130
                        135
<210> 64
<211> 1656
<212> DNA
<213> Saccharomyces cerevisiae
<400> 64
atgcgttggt ctgtccttga tacagtgcta ttgaccgtga tttcctttca tctaatccaa 60
gctccattca ccaaggtgga agaggtttt aatattcaag ccattcatga tattttaacc 120
tacagcgtat ttgatatctc ccaatatgac cacttgaaat ttcctggagt agtccctaga 180
acattegttg gtgetgtgat tattgeaatg etttegagae ettatettta ettgagttet 240
ttgatccaaa cttccaggcc tacgtctata gatgttcaat tggtcgttag ggggattgtt 300
ggcctcacca atgggctttc ttttatctat ttaaaqaatt gtttqcaaqa tatgtttqat 360
gaaatcactg aaaagaaaaa ggaaqaaaat gaaqacaaqq atatatacat ttacqataqc 420
gctggtacat ggtttctttt atttttaatt ggcagtttcc acctcatgtt ctacagcact 480
aggactetge etaattttgt catgactetg cetetaacea aegtegeatt ggggtgggtt 540
ttattgggtc gttataatgc agctatattc ctatctgcgc tcgtggcaat tgtatttaga 600
ctggaagtgt cagctctcag tgctggtatt gctctattta gcgtcatctt caagaagatt 660
tctttattcg atgctatcaa attcggtatc tttggcttgg gacttggttc cgccatcagt 720
atcaccgttg attcatattt ctggcaagaa tggtgtctac ctgaggtaga tggtttcttg 780
ttcaacgtgg ttgcgggtta cgcttccaag tggggtgtgg agccagttac tgcttatttc 840
acgcattact tgagaatgat gtttatgcca ccaactgttt tactattgaa ttacttcggc 900
tataaattag cacctgcaaa attaaaaatt gtctcactag catctctttt ccacattatc 960
gtettateet tteaacetea caaagaatgg agatteatea tetaegetgt tecatetate 1020
atgttgctag gtgccacagg agcagcacat ctatgggaga atatgaaagt aaaaaagatt 1080
atggcgttct tgtatatatc aagaatgaat tatccaggcg gcgaggcttt aacttctttt 1200
aatgacatga ttgtggaaaa aaatattaca aacgctacag ttcatatcag catacctcct 1260
tgcatgacag gtgtcacttt atttggtgaa ttgaactacg gtgtgtacgg catcaattac 1320
gataagactg aaaatacgac tttactgcag gaaatgtggc cctcctttga tttcttgatc 1380
acccacgage caaccgcete teaattgeea ttegagaata agactaceaa ceattgggag 1440
ctagttaaca caacaaagat gtttactgga tttgacccaa cctacattaa gaactttgtt 1500
ttccaagaga gagtgaatgt tttgtctcta ctcaaacaga tcattttcga caagacccct 1560
accgtttttt tgaaagaatt gacggccaat tcgattgtta aaagcgatgt cttcttcacc 1620
tataagagaa tcaaacaaga tgaaaaaact gattga
<210> 65
<211> 551
<212> PRT
<213> Saccharomyces cerevisiae
<400> 65
```

Met Arg Trp Ser Val Leu Asp Thr Val Leu Leu Thr Val Ile Ser Phe

```
His Leu Ile Gln Ala Pro Phe Thr Lys Val Glu Glu Ser Phe Asn Ile
                                25
Gln Ala Ile His Asp Ile Leu Thr Tyr Ser Val Phe Asp Ile Ser Gln
Tyr Asp His Leu Lys Phe Pro Gly Val Val Pro Arg Thr Phe Val Gly
                       55
Ala Val Ile Ile Ala Met Leu Ser Arg Pro Tyr Leu Tyr Leu Ser Ser
                    70
                                        75
Leu Ile Gln Thr Ser Arg Pro Thr Ser Ile Asp Val Gln Leu Val Val
                                    90
Arg Gly Ile Val Gly Leu Thr Asn Gly Leu Ser Phe Ile Tyr Leu Lys
                                105
Asn Cys Leu Gln Asp Met Phe Asp Glu Ile Thr Glu Lys Lys Glu
                            120
Glu Asn Glu Asp Lys Asp Ile Tyr Ile Tyr Asp Ser Ala Gly Thr Trp
                        135
                                            140
Phe Leu Phe Leu Ile Gly Ser Phe His Leu Met Phe Tyr Ser Thr
                    150
                                        155
Arg Thr Leu Pro Asn Phe Val Met Thr Leu Pro Leu Thr Asn Val Ala
                165
                                   170
Leu Gly Trp Val Leu Leu Gly Arg Tyr Asn Ala Ala Ile Phe Leu Ser
           180
                                185
Ala Leu Val Ala Ile Val Phe Arg Leu Glu Val Ser Ala Leu Ser Ala
                           200
                                                205
Gly Ile Ala Leu Phe Ser Val Ile Phe Lys Lys Ile Ser Leu Phe Asp
                       215
Ala Ile Lys Phe Gly Ile Phe Gly Leu Gly Leu Gly Ser Ala Ile Ser
                  230
                                        235
Ile Thr Val Asp Ser Tyr Phe Trp Gln Glu Trp Cys Leu Pro Glu Val
                                   250
Asp Gly Phe Leu Phe Asn Val Val Ala Gly Tyr Ala Ser Lys Trp Gly
                               265
Val Glu Pro Val Thr Ala Tyr Phe Thr His Tyr Leu Arg Met Met Phe
                           280
Met Pro Pro Thr Val Leu Leu Asn Tyr Phe Gly Tyr Lys Leu Ala
                       295
                                            300
Pro Ala Lys Leu Lys Ile Val Ser Leu Ala Ser Leu Phe His Ile Ile
                   310
                                        315
Val Leu Ser Phe Gln Pro His Lys Glu Trp Arg Phe Ile Ile Tyr Ala
               325
                                    330
Val Pro Ser Ile Met Leu Leu Gly Ala Thr Gly Ala Ala His Leu Trp
                               345
Glu Asn Met Lys Val Lys Lys Ile Thr Asn Val Leu Cys Leu Ala Ile
                           360
Leu Pro Leu Ser Ile Met Thr Ser Phe Phe Ile Ser Met Ala Phe Leu
                       375
                                            380
Tyr Ile Ser Arg Met Asn Tyr Pro Gly Gly Glu Ala Leu Thr Ser Phe
                   390
                                        395
Asn Asp Met Ile Val Glu Lys Asn Ile Thr Asn Ala Thr Val His Ile
               405
                                   410
Ser Ile Pro Pro Cys Met Thr Gly Val Thr Leu Phe Gly Glu Leu Asn
           420
                               425
                                                   430
Tyr Gly Val Tyr Gly Ile Asn Tyr Asp Lys Thr Glu Asn Thr Thr Leu
                           440
                                               445
Leu Gln Glu Met Trp Pro Ser Phe Asp Phe Leu Ile Thr His Glu Pro
                       455
                                           460
Thr Ala Ser Gln Leu Pro Phe Glu Asn Lys Thr Thr Asn His Trp Glu
```

```
470
                                         475
Leu Val Asn Thr Thr Lys Met Phe Thr Gly Phe Asp Pro Thr Tyr Ile
                485
                                     490
Lys Asn Phe Val Phe Gln Glu Arg Val Asn Val Leu Ser Leu Leu Lys
                                 505
Gln Ile Ile Phe Asp Lys Thr Pro Thr Val Phe Leu Lys Glu Leu Thr
                             520
                                                 525
Ala Asn Ser Ile Val Lys Ser Asp Val Phe Phe Thr Tyr Lys Arg Ile
                        535
Lys Gln Asp Glu Lys Thr Asp
545
<210> 66
<211> 840
<212> DNA
<213> Pichia pastoris
<400> 66
teggtegaga atgataactg aagaacteaa aateteteac aettteateg ttaetgtaet 60
ggcaatcatt gcatttcagc ctcataaaga atggagattt atagtttaca ttgttccacc 120
acttgtcatc accatatcta cagtacttgc acaactaccc aggagattca caatcgtcaa 180
agttgctgtt tttctcctaa gtttcggctc tttgctcata tccctgtcgt ttcttttcat 240
ctcatcgtat aactaccctg ggggtgaagc tttacagcat ttgaacgaga aactccttct 300
actggaccaa agttccctac ctgttgatat taaggttcat atggatgtcc ctgcatgcat 360
gactggggtg actttatttg gttacttgga taactcaaaa ttgaacaatt taagaattgt 420
ctatgataaa acagaagacg agtcgctgga cacaatctgg gattctttca attatgtcat 480
ctccgaaatt gacttggatt cttcgactgc tcccaaatgg gagggggatt ggctgaagat 540
tgatgttgtc caaggctaca acggcatcaa taaacaatct atcaaaaata caattttcaa 600
ttatggaata cttaaacgga tgataagaga cgcaaccaaa cttgatgttg gatttattcg 660
tacggtcttt cgatccttca taaaatttga tgataaatta ttcatttatg agaggagcag 720
tcaaacctga aaatatatac ctcatttgtt caatttggtg taaagagtgt ggcggataga 780
cttcttgtaa atcaggaaag ctacaattcc aattgctgca aaaaatacca atgcccataa 840
<210> 67
<211> 239
<212> PRT
<213> Pichia pastoris
<400> 67
Arg Met Ile Thr Glu Glu Leu Lys Ile Ser His Thr Phe Ile Val Thr
                                    10
Val Leu Ala Ile Ile Ala Phe Gln Pro His Lys Glu Trp Arg Phe Ile
                                25
Val Tyr Ile Val Pro Pro Leu Val Ile Thr Ile Ser Thr Val Leu Ala
                            40
Gln Leu Pro Arg Arg Phe Thr Ile Val Lys Val Ala Val Phe Leu Leu
Ser Phe Gly Ser Leu Leu Ile Ser Leu Ser Phe Leu Phe Ile Ser Ser
                                        75
Tyr Asn Tyr Pro Gly Gly Glu Ala Leu Gln His Leu Asn Glu Lys Leu
                                    90
Leu Leu Leu Asp Gln Ser Ser Leu Pro Val Asp Ile Lys Val His Met
            100
                                105
                                                     110
Asp Val Pro Ala Cys Met Thr Gly Val Thr Leu Phe Gly Tyr Leu Asp
                            120
                                                125
Asn Ser Lys Leu Asn Asn Leu Arg Ile Val Tyr Asp Lys Thr Glu Asp
```

```
135
Glu Ser Leu Asp Thr Ile Trp Asp Ser Phe Asn Tyr Val Ile Ser Glu
                  150
                                      155
Ile Asp Leu Asp Ser Ser Thr Ala Pro Lys Trp Glu Gly Asp Trp Leu
               165
                                   170
Lys Ile Asp Val Val Gln Gly Tyr Asn Gly Ile Asn Lys Gln Ser Ile
                               185
Lys Asn Thr Ile Phe Asn Tyr Gly Ile Leu Lys Arg Met Ile Arg Asp
                           200
       195
Ala Thr Lys Leu Asp Val Gly Phe Ile Arg Thr Val Phe Arg Ser Phe
                       215
Ile Lys Phe Asp Asp Lys Leu Phe Ile Tyr Glu Arg Ser Ser Gln
                   230
<210> 68
<211> 239
<212> PRT
<213> Pichia pastoris
<220>
<221> MOD RES
<222> (62)...(80)
<223> Xaa is a variable amino acid
<400> 68
Arg Met Ile Thr Glu Glu Leu Lys Ile Ser His Thr Phe Ile Val Thr
               - 5
                                   10
Val Leu Ala Ile Ile Ala Phe Gln Pro His Lys Glu Trp Arg Phe Ile
                               25
Val Tyr Ile Val Pro Pro Leu Val Ile Thr Ile Ser Thr Val Leu Ala
                           40
Gln Leu Pro Arg Arg Phe Thr Ile Val Lys Val Ala Val Xaa Xaa Xaa
75
Tyr Asn Tyr Pro Gly Gly Glu Ala Leu Gln His Leu Asn Glu Lys Leu
                                   90
Leu Leu Leu Asp Gln Ser Ser Leu Pro Val Asp Ile Lys Val His Met
                              105
Asp Val Pro Ala Cys Met Thr Gly Val Thr Leu Phe Gly Tyr Leu Asp
                           120
                                              125
Asn Ser Lys Leu Asn Asn Leu Arg Ile Val Tyr Asp Lys Thr Glu Asp
                       135
Glu Ser Leu Asp Thr Ile Trp Asp Ser Phe Asn Tyr Val Ile Ser Glu
                  150
                                      155
Ile Asp Leu Asp Ser Ser Thr Ala Pro Lys Trp Glu Gly Asp Trp Leu
               165
                                  170
Lys Ile Asp Val Val Gln Gly Tyr Asn Gly Ile Asn Lys Gln Ser Ile
                               185
Lys Asn Thr Ile Phe Asn Tyr Gly Ile Leu Lys Arg Met Ile Arg Asp
                           200
                                              205
Ala Thr Lys Leu Asp Val Gly Phe Ile Arg Thr Val Phe Arg Ser Phe
                      215
                                          220
Ile Lys Phe Asp Asp Lys Leu Phe Ile Tyr Glu Arg Ser Ser Gln
```

```
<210> 69
 <211> 245
 <212> PRT
 <213> Saccharomyces cerevisiae
 <400> 69
 Lys Leu Ala Pro Ala Lys Leu Lys Ile Val Ser Leu Ala Ser Leu Phe
 1
 His Ile Ile Val Leu Ser Phe Gln Pro His Lys Glu Trp Arg Phe Ile
 Ile Tyr Ala Val Pro Ser Ile Met Leu Leu Gly Ala Thr Gly Ala Ala
 His Leu Trp Glu Asn Met Lys Val Lys Lys Ile Thr Asn Val Leu Cys
                        55
 Leu Ala Ile Leu Pro Leu Ser Ile Met Thr Ser Phe Phe Ile Ser Met
                                        75
 Ala Phe Leu Tyr Ile Ser Arg Met Asn Tyr Pro Gly Gly Glu Ala Leu
                                    90
 Thr Ser Phe Asn Asp Met Ile Val Glu Lys Asn Ile Thr Asn Ala Thr
            100
                                105
 Val His Ile Ser Ile Pro Pro Cys Met Thr Gly Val Thr Leu Phe Gly
                            120
                                               125
 Glu Leu Asn Tyr Gly Val Tyr Gly Ile Asn Tyr Asp Lys Thr Glu Asn
                        135
                                           140
 Thr Thr Leu Leu Gln Glu Met Trp Pro Ser Phe Asp Phe Leu Ile Thr
                   150
                                       155
 His Glu Pro Thr Ala Ser Gln Leu Pro Phe Glu Asn Lys Thr Thr Asn
               1.65
                                   170
 His Trp Glu Leu Val Asn Thr Thr Lys Met Phe Thr Gly Phe Asp Pro
            180
                               185
 Thr Tyr Ile Lys Asn Phe Val Phe Gln Glu Arg Val Asn Val Leu Ser
                           200
 Leu Leu Lys Gln Ile Ile Phe Asp Lys Thr Pro Thr Val Phe Leu Lys
                       215
                                           220
 Glu Leu Thr Ala Asn Ser Ile Val Lys Ser Asp Val Phe Phe Thr Tyr
                                       235
 Lys Arg Ile Lys Gln
. <210> 70
 <211> 141
 <212> PRT
 <213> Pichia pastoris
 <220>
 <221> MOD RES
 <222> (43)...(61)
 <223> Xaa is a variable amino acid
 <400> 70
 Ile Ile Ala Phe Gln Pro His Lys Glu Trp Arg Phe Ile Val Tyr Ile
                                   10
 Val Pro Pro Leu Val Ile Thr Ile Ser Thr Val Leu Ala Gln Leu Pro
                               25
Arg Arg Phe Thr Ile Val Lys Val Ala Val Xaa Xaa Xaa Xaa Xaa
                           40
                                               4.5
```

```
Pro Gly Gly Glu Ala Leu Gln His Leu Asn Glu Lys Leu Leu Leu
                   70
                                      75
Asp Gln Ser Ser Leu Pro Val Asp Ile Lys Val His Met Asp Val Pro
               85
                                   90
Ala Cys Met Thr Gly Val Thr Leu Phe Gly Tyr Leu Asp Asn Ser Lys
           100
                               105
Leu Asn Asn Leu Arg Ile Val Tyr Asp Lys Thr Glu Asp Glu Ser Leu
                           120
Asp Thr Ile Trp Asp Ser Phe Asn Tyr Val Ile Ser Glu
                       135
<210> 71
<211> 137
<212> PRT
<213> Schizosaccharomyces pombe
<400> 71
Val Tyr Ser Phe Leu Gly His Lys Glu Trp Arg Phe Ile Ile Tyr Ser
1
                                                      1.5
                                  10
Ile Pro Trp Phe Asn Ala Ala Ser Ala Ile Gly Ala Ser Leu Cys Phe
                              25
Asn Ala Ser Lys Phe Gly Lys Lys Ile Phe Glu Ile Leu Arg Leu Met
                          40
Phe Phe Ser Gly Ile Ile Phe Gly Phe Ile Gly Ser Ser Phe Leu Leu
                                          60
Tyr Val Phe Gln Tyr Ala Tyr Pro Gly Gly Leu Ala Leu Thr Arg Leu
                  70
                                      75
Tyr Glu Ile Glu Asn His Pro Gln Val Ser Val His Met Asp Val Tyr
Pro Cys Met Thr Gly Ile Thr Arg Phe Ser Gln Leu Pro Ser Trp Tyr
                              105
Tyr Asp Lys Thr Glu Asp Pro Lys Met Leu Ser Asn Ser Leu Phe Ile
                          120
Ser Gln Phe Asp Tyr Leu Ile Thr Glu
                       135
<210> 72
<211> 143
<212> PRT
<213> Pichia pastoris
<220>
<221> MOD RES
<222> (45)...(63)
<223> Xaa is a variable amino acid
Leu Ala Ile Ile Ala Phe Gln Pro His Lys Glu Trp Arg Phe Ile Val
                                  10
Tyr Ile Val Pro Pro Leu Val Ile Thr Ile Ser Thr Val Leu Ala Gln
           20
                              25
Leu Pro Arg Arg Phe Thr Ile Val Lys Val Ala Val Xaa Xaa Xaa Xaa
                          40
                                              4.5
```

```
Asn Tyr Pro Gly Gly Glu Ala Leu Gln His Leu Asn Glu Lys Leu Leu
Leu Leu Asp Gln Ser Ser Leu Pro Val Asp Ile Lys Val His Met Asp
                                     90
Val Pro Ala Cys Met Thr Gly Val Thr Leu Phe Gly Tyr Leu Asp Asn
            100
                                105
Ser Lys Leu Asn Asn Leu Arg Ile Val Tyr Asp Lys Thr Glu Asp Glu
                            120
Ser Leu Asp Thr Ile Trp Asp Ser Phe Asn Tyr Val Ile Ser Glu
    130
                        135
<210> 73
<211> 137
<212> PRT
<213> Homo sapiens
<400> 73
Met Ala Leu Tyr Ser Leu Leu Pro His Lys Glu Leu Arg Phe Ile Ile
                                     10
Tyr Ala Phe Pro Met Leu Asn Ile Thr Ala Ala Arg Gly Cys Ser Tyr
            20
                                25
                                                     30
Leu Leu Asn Asn Tyr Lys Lys Ser Trp Leu Tyr Lys Ala Gly Ser Leu
                            40
                                                 45
Leu Val Ile Gly His Leu Val Val Asn Ala Ala Tyr Ser Ala Thr Ala
                        55
                                             60
Leu Tyr Val Ser His Phe Asn Tyr Pro Gly Gly Val Ala Met Gln Arg
                    70
                                        75
Leu His Gln Leu Val Pro Pro Gln Thr Asp Val Leu Leu His Ile Asp
                85
                                    90
Val Ala Ala Gln Thr Gly Val Ser Arg Phe Leu Gln Val Asn Ser
                                105
Ala Trp Arg Tyr Asp Lys Arg Glu Asp Val Gln Pro Gly Thr Gly Met
                            120
Leu Ala Tyr Thr His Ile Leu Met Glu
                        135
<210> 74
<211> 1635
<212> DNA
<213> Saccharomyces cerevisiae
<400> 74
atggccattg gcaaaaggtt actggtgaac aaaccagcag aagaatcatt ttatgcttct 60
ccaatgtatg attttttgta tccgtttagg ccagtgggga accaatggct gccagaatat 120
attatetttg tatgtgetgt aataetgagg tgeacaattg gaettggtee atattetggg 180
aaaggcagtc caccgctgta cggcgatttt gaggctcaga gacattggat ggaaattacg 240
caacatttac cgctttctaa gtggtactgg tatgatttgc aatactgggg attggactat 300
ccaccattaa cagcatttca ttcgtacctt ctgggcctaa ttggatcttt tttcaatcca 360
tcttggtttg cactagaaaa gtcacgtggc tttgaatccc ccgataatgg cctgaaaaca 420
tatatgcgtt ctactgtcat cattagcgac atattgtttt actttcctgc agtaatatac 480
tttactaagt ggcttggtag atatcgaaac cagtcgccca taggacaatc tattgcggca 540
tcagcgattt tgttccaacc ttcattaatg ctcattgacc atgggcactt tcaatataat 600
tcagtcatgc ttggccttac tgcttatgcc ataaataact tattagatga gtattatgct 660
atggcggccg tttgttttgt cctatccatt tgttttaaac aaatggcatt gtattatgca 720
ecgatttttt ttgettatet attaagtega teattgetgt teeceaaatt taacataget 780
agattgacgg ttattgcgtt tgcaacactc gcaacttttg ctataatatt tgcgccatta 840
```

```
tatttcttgg gaggaggatt aaagaatatt caccaatgta ttcacaggat attccctttt 900
gccaggggca tcttcgaaga caaggttgct aacttctggt gcgttacgaa cgtgtttgta 960
aaatacaagg aaagattcac tatacaacaa ctccagctat attcattgat tgccaccgtg 1020
attggtttct taccagccat gataatgaca ttacttcatc ccaaaaaagca tcttctccca 1080
tacgtgttaa tcgcatgttc gatgtccttt tttcttttta gctttcaagt acatgagaaa 1140
actatectea teccaetttt geetattaea etaetetaet eetetaetga ttggaatgtt 1200
ctatctcttg taagttggat aaacaatgtg gctttgttta cgctatggcc tttgttgaaa 1260
aaggacggtc ttcatttaca gtatgccgta tctttcttac taagcaattg gctgattgga 1320
aatttcagtt ttattacacc aaggttcttg ccaaaatctt taactcctgg cccttctatc 1380
agcagcatca atagcgacta tagaagaaga agcttactgc catataatgt ggtttggaaa 1440
agttttatca taggaacgta tattgctatg ggcttttatc atttcttaga tcaatttgta 1500
gcacctccat cgaaatatcc agacttgtgg gtgttgttga actgtgctgt tgggttcatt 1560
tgctttagca tattttggct atggtcttat tacaagatat tcacttccgg tagcaaatcc 1620
atgaaggact tgtag
<210> 75
<211> 544
<212> PRT
<213> Saccharomyces cerevisiae
<400> 75
Met Ala Ile Gly Lys Arg Leu Leu Val Asn Lys Pro Ala Glu Glu Ser
 1
                                    10
Phe Tyr Ala Ser Pro Met Tyr Asp Phe Leu Tyr Pro Phe Arg Pro Val
            20
                                25
Gly Asn Gln Trp Leu Pro Glu Tyr Ile Ile Phe Val Cys Ala Val Ile
                            40
                                                 45
Leu Arg Cys Thr Ile Gly Leu Gly Pro Tyr Ser Gly Lys Gly Ser Pro
                        55
Pro Leu Tyr Gly Asp Phe Glu Ala Gln Arg His Trp Met Glu Ile Thr
                    70
                                        75
Gln His Leu Pro Leu Ser Lys Trp Tyr Trp Tyr Asp Leu Gln Tyr Trp
                                    90
Gly Leu Asp Tyr Pro Pro Leu Thr Ala Phe His Ser Tyr Leu Leu Gly
                                105
Leu Ile Gly Ser Phe Phe Asn Pro Ser Trp Phe Ala Leu Glu Lys Ser
                            120
Arg Gly Phe Glu Ser Pro Asp Asn Gly Leu Lys Thr Tyr Met Arg Ser
                        135
                                            140
Thr Val Ile Ile Ser Asp Ile Leu Phe Tyr Phe Pro Ala Val Ile Tyr
                    150
                                        155
Phe Thr Lys Trp Leu Gly Arg Tyr Arg Asn Gln Ser Pro Ile Gly Gln
                165
                                    170
Ser Ile Ala Ala Ser Ala Ile Leu Phe Gln Pro Ser Leu Met Leu Ile
            180
                                185
                                                     190
Asp His Gly His Phe Gln Tyr Asn Ser Val Met Leu Gly Leu Thr Ala
        195
                            200
                                                 205
Tyr Ala Ile Asn Asn Leu Leu Asp Glu Tyr Tyr Ala Met Ala Ala Val
                        215
                                            220
Cys Phe Val Leu Ser Ile Cys Phe Lys Gln Met Ala Leu Tyr Tyr Ala
                    230
                                        235
Pro Ile Phe Phe Ala Tyr Leu Leu Ser Arg Ser Leu Leu Phe Pro Lys
                                    250
Phe Asn Ile Ala Arg Leu Thr Val Ile Ala Phe Ala Thr Leu Ala Thr
                                265
Phe Ala Ile Ile Phe Ala Pro Leu Tyr Phe Leu Gly Gly Leu Lys
                            280
                                                285
Asn Ile His Gln Cys Ile His Arg Ile Phe Pro Phe Ala Arg Gly Ile
```

```
295
Phe Glu Asp Lys Val Ala Asn Phe Trp Cys Val Thr Asn Val Phe Val
                    310
                                         315
Lys Tyr Lys Glu Arg Phe Thr Ile Gln Gln Leu Gln Leu Tyr Ser Leu
                325
                                     330
Ile Ala Thr Val Ile Gly Phe Leu Pro Ala Met Ile Met Thr Leu Leu
                                 345
His Pro Lys Lys His Leu Leu Pro Tyr Val Leu Ile Ala Cys Ser Met
        355
                             360
Ser Phe Phe Leu Phe Ser Phe Gln Val His Glu Lys Thr Ile Leu Ile
                        375
Pro Leu Leu Pro Ile Thr Leu Leu Tyr Ser Ser Thr Asp Trp Asn Val
385
Leu Ser Leu Val Ser Trp Ile Asn Asn Val Ala Leu Phe Thr Leu Trp
                                     410
Pro Leu Leu Lys Lys Asp Gly Leu His Leu Gln Tyr Ala Val Ser Phe
            420
                                 425
                                                      430
Leu Leu Ser Asn Trp Leu Ile Gly Asn Phe Ser Phe Ile Thr Pro Arg
        435
                             440
Phe Leu Pro Lys Ser Leu Thr Pro Gly Pro Ser Ile Ser Ser Ile Asn
    450
                        455
                                             460
Ser Asp Tyr Arg Arg Arg Ser Leu Leu Pro Tyr Asn Val Val Trp Lys
465
                    470
                                         475
                                                              480
Ser Phe Ile Ile Gly Thr Tyr Ile Ala Met Gly Phe Tyr His Phe Leu
                485
                                     490
Asp Gln Phe Val Ala Pro Pro Ser Lys Tyr Pro Asp Leu Trp Val Leu
            500
                                505
                                                     510
Leu Asn Cys Ala Val Gly Phe Ile Cys Phe Ser Ile Phe Trp Leu Trp
        515
                            520
Ser Tyr Tyr Lys Ile Phe Thr Ser Gly Ser Lys Ser Met Lys Asp Leu
    530
                        535
```

<210> 76

<211> 1644

<212> DNA

<213> Pichia pastoris

<400> 76

atgccacata aaagaacgcc ctctagcagt ctgctgtatg caagaattcc agggatctct 60 tttgaaaact ctccggtgtt tgattttttg tctccttttg gacccgctcc taatcaatgg 120 tectatteeg getteaacae ecetecaatg tatggggatt ttgaagetea gaggeattgg 240 atggaaatta ctcagcattt atccatagaa aaatggtact tctacgactt gcaatattgg 300 gggcttgact atcctccctt gacagccttt cattcatact tctttggcaa attaggcagc 360 ttcatcaatc cagcatggtt tgctttagac gtctccagag ggtttgaatc agtggatcta 420 aaatcgtaca tgagggcgac cgcaattctc agtgagctgt tatgttttat tccagctgtc 480 atttggtatt gtcgttggat gggacttaac tacttcaatc aaaacgccat tgagcaaact 540 ataatagcgt ctgctattct tttcaatcca tctttaatta tcatagatca tggccacttc 600 cagtacaact cagttatgct aggttttgct ttattatcca tattaaatct gttgtacgat 660 aattttgcat tagcggctat ttttttcgtt ctttcaataa gctttaagca aatggctctc 720 tattatagcc ccatcatgtt tttttacatg ctgagtgtga gttgttggcc tttgaaaaac 780 ttcaacttgt tgagattggc tactatcagt attgcagtac tcttgacttt tgcaactcta 840 ttactgcctt ttgtattagt agatgggatg tcacaaattg gccaaatatt attcagagtt 900 ttcccgtttt caagaggctt gtttgaggat aaggtggcca acttttggtg tacaacgaat 960 atactggtaa agtacaaaca gttattcact gacaaaaccc ttactaggat atcgctagta 1020 gcaactttga ttgcaattag tccgtcttgc ttcatcattt ttactcaccc aaagaaggtt 1080 ttactaccgt gggcttttgc tgcttgctct tgggcgttct atcttttctc tttccaagtc 1140

```
cacgagaaat cagttttagt tccattgatg cctaccactc tattactggt agaaaaagac 1200
ttggacatca tctcaatggt ctgctggatt tctaatattg ccttcttcag catgtggcct 1260
ctattaaaaa gagacgggct ggctttggaa tattttgtct tgggaatatt gagtaattgg 1320
ctgattggaa acctcaattg gattagtaaa tggcttgtcc ccagtttcct gattccaggg 1380
cctactctct ccaaaaaagt tcctaaaaga gatactaaaa cagttgttca tactcactgg 1440
ttttgggggt cagtaacatt cgtttcatac ctcggagcta cagttatcca gttcgtagat 1500
tggctgtacc ttccacctgc caagtatcca gatttgtggg ttattttgaa cactacattg 1560
tegtttgett gtttegggtt gttttggeta tggattaact acaatetgta cattttgegt 1620
gattttaagc ttaaagatgc ttag
<210> 77
<211> 547
<212> PRT
<213> Pichia pastoris
<400> 77
Met Pro His Lys Arg Thr Pro Ser Ser Ser Leu Leu Tyr Ala Arg Ile
Pro Gly Ile Ser Phe Glu Asn Ser Pro Val Phe Asp Phe Leu Ser Pro
                                25
Phe Gly Pro Ala Pro Asn Gln Trp Val Ala Arg Tyr Ile Ile Ile Ile
                            40
Phe Ala Ile Leu Ile Arg Leu Ala Val Gly Leu Gly Ser Tyr Ser Gly
                        55
                                            60
Phe Asn Thr Pro Pro Met Tyr Gly Asp Phe Glu Ala Gln Arg His Trp
                    70
                                        75
Met Glu Ile Thr Gln His Leu Ser Ile Glu Lys Trp Tyr Phe Tyr Asp
               85
                                    90
                                                        95
Leu Gln Tyr Trp Gly Leu Asp Tyr Pro Pro Leu Thr Ala Phe His Ser
            100
                                105
                                                    110
Tyr Phe Phe Gly Lys Leu Gly Ser Phe Ile Asn Pro Ala Trp Phe Ala
       115
                            120
                                                125
Leu Asp Val Ser Arg Gly Phe Glu Ser Val Asp Leu Lys Ser Tyr Met
                        135
                                            140
Arg Ala Thr Ala Ile Leu Ser Glu Leu Cys Phe Ile Pro Ala Val
                    150
                                        155
Ile Trp Tyr Cys Arg Trp Met Gly Leu Asn Tyr Phe Asn Gln Asn Ala
               165
                                    170
Ile Glu Gln Thr Ile Ile Ala Ser Ala Ile Leu Phe Asn Pro Ser Leu
           180 ·
                                185
                                                    190
Ile Ile Ile Asp His Gly His Phe Gln Tyr Asn Ser Val Met Leu Gly
                            200
                                                205
Phe Ala Leu Leu Ser Ile Leu Asn Leu Leu Tyr Asp Asn Phe Ala Leu
                        215
                                            220
Ala Ala Ile Phe Phe Val Leu Ser Ile Ser Phe Lys Gln Met Ala Leu
                    230
                                        235
Tyr Tyr Ser Pro Ile Met Phe Phe Tyr Met Leu Ser Val Ser Cys Trp
               245
                                    250
Pro Leu Lys Asn Phe Asn Leu Leu Arg Leu Ala Thr Ile Ser Ile Ala
            260
                                265
Val Leu Leu Thr Phe Ala Thr Leu Leu Pro Phe Val Leu Val Asp
                            280
                                                285
Gly Met Ser Gln Ile Gly Gln Ile Leu Phe Arg Val Phe Pro Phe Ser
                       295
Arg Gly Leu Phe Glu Asp Lys Val Ala Asn Phe Trp Cys Thr Thr Asn
                   310
                                        315
Ile Leu Val Lys Tyr Lys Gln Leu Phe Thr Asp Lys Thr Leu Thr Arg
               325
                                    330
                                                        335
```

```
Ile Ser Leu Val Ala Thr Leu Ile Ala Ile Ser Pro Ser Cys Phe Ile
                                345
Ile Phe Thr His Pro Lys Lys Val Leu Pro Trp Ala Phe Ala Ala
                            360
Cys Ser Trp Ala Phe Tyr Leu Phe Ser Phe Gln Val His Glu Lys Ser
                        375
                                            380
Val Leu Val Pro Leu Met Pro Thr Thr Leu Leu Leu Val Glu Lys Asp
                    390
                                        395
Leu Asp Ile Ile Ser Met Val Cys Trp Ile Ser Asn Ile Ala Phe Phe
                405
                                    410
Ser Met Trp Pro Leu Leu Lys Arg Asp Gly Leu Ala Leu Glu Tyr Phe
                                425
Val Leu Gly Ile Leu Ser Asn Trp Leu Ile Gly Asn Leu Asn Trp Ile
        435
                            440
Ser Lys Trp Leu Val Pro Ser Phe Leu Ile Pro Gly Pro Thr Leu Ser
                        455
                                            460
Lys Lys Val Pro Lys Arg Asp Thr Lys Thr Val Val His Thr His Trp
                    470
                                        475
Phe Trp Gly Ser Val Thr Phe Val Ser Tyr Leu Gly Ala Thr Val Ile
                485
                                    490
Gln Phe Val Asp Trp Leu Tyr Leu Pro Pro Ala Lys Tyr Pro Asp Leu
                                505
                                                    510
Trp Val Ile Leu Asn Thr Thr Leu Ser Phe Ala Cys Phe Gly Leu Phe
                           520
                                                525
Trp Leu Trp Ile Asn Tyr Asn Leu Tyr Ile Leu Arg Asp Phe Lys Leu
   530
                       535
Lys Asp Ala
545
<210> 78
<211> 527
<212> PRT
<213> Pichia pastoris
<220>
<221> MOD RES
<222> (23)...(37)
<223> Xaa is a variable amino acid
<220>
<221> MOD RES
<222> (366)...(378)
<223> Xaa is a variable amino acid
<400> 78
Ser Phe Glu Asn Ser Pro Val Phe Asp Phe Leu Ser Pro Phe Gly Pro
                5
                                    10
Ala Pro Asn Gln Trp Val Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa
           20
                                25
Xaa Xaa Xaa Xaa Val Gly Leu Gly Ser Tyr Ser Gly Phe Asn Thr
                                                45
Pro Pro Met Tyr Gly Asp Phe Glu Ala Gln Arg His Trp Met Glu Ile
                       55
Thr Gln His Leu Ser Ile Glu Lys Trp Tyr Phe Tyr Asp Leu Gln Tyr
                   70
                                        75
Trp Gly Leu Asp Tyr Pro Pro Leu Thr Ala Phe His Ser Tyr Phe Phe
```

```
Gly Lys Leu Gly Ser Phe Ile Asn Pro Ala Trp Phe Ala Leu Asp Val
                               105
Ser Arg Gly Phe Glu Ser Val Asp Leu Lys Ser Tyr Met Arg Ala Thr
        115
                           120
Ala Ile Leu Ser Glu Leu Leu Cys Phe Ile Pro Ala Val Ile Trp Tyr
                       135
                                           140
Cys Arg Trp Met Gly Leu Asn Tyr Phe Asn Gln Asn Ala Ile Glu Gln
                   150
                                       155
Thr Ile Ile Ala Ser Ala Ile Leu Phe Asn Pro Ser Leu Ile Ile Ile
               165
                                   170
Asp His Gly His Phe Gln Tyr Asn Ser Val Met Leu Gly Phe Ala Leu
           180
                               185
Leu Ser Ile Leu Asn Leu Leu Tyr Asp Asn Phe Ala Leu Ala Ile
                           200
Phe Phe Val Leu Ser Ile Ser Phe Lys Gln Met Ala Leu Tyr Tyr Ser
                       215
Pro Ile Met Phe Phe Tyr Met Leu Ser Val Ser Cys Trp Pro Leu Lys
                   230
                                       235
Asn Phe Asn Leu Leu Arg Leu Ala Thr Ile Ser Ile Ala Val Leu Leu
                                   250
               245
Thr Phe Ala Thr Leu Leu Pro Phe Val Leu Val Asp Gly Met Ser
                               265
Gln Ile Gly Gln Ile Leu Phe Arg Val Phe Pro Phe Ser Arg Gly Leu
                          280
       275
                                               285
Phe Glu Asp Lys Val Ala Asn Phe Trp Cys Thr Thr Asn Ile Leu Val
                       295
                                           300
Lys Tyr Lys Gln Leu Phe Thr Asp Lys Thr Leu Thr Arg Ile Ser Leu
                   310
                                       315
Val Ala Thr Leu Ile Ala Ile Ser Pro Ser Cys Phe Ile Ile Phe Thr
              325
                                   330
His Pro Lys Lys Val Leu Leu Pro Trp Ala Phe Ala Ala Cys Ser Trp
           340
                               345
Ala Phe Tyr Leu Phe Ser Phe Gln Val His Glu Lys Ser Xaa Xaa
                           360
Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Glu Lys Asp Leu Asp Ile
                       375
                                           380
Ile Ser Met Val Cys Trp Ile Ser Asn Ile Ala Phe Phe Ser Met Trp
                   390
                                       395
Pro Leu Leu Lys Arg Asp Gly Leu Ala Leu Glu Tyr Phe Val Leu Gly
               405
                                   410
Ile Leu Ser Asn Trp Leu Ile Gly Asn Leu Asn Trp Ile Ser Lys Trp
           420
                               425
Leu Val Pro Ser Phe Leu Ile Pro Gly Pro Thr Leu Ser Lys Lys Val
                           440
                                               445
Pro Lys Arg Asp Thr Lys Thr Val Val His Thr His Trp Phe Trp Gly
                      455
                                           460
Ser Val Thr Phe Val Ser Tyr Leu Gly Ala Thr Val Ile Gln Phe Val
                  470
                                       475
Asp Trp Leu Tyr Leu Pro Pro Ala Lys Tyr Pro Asp Leu Trp Val Ile
               485
                                   490
Leu Asn Thr Thr Leu Ser Phe Ala Cys Phe Gly Leu Phe Trp Leu Trp
           500
                               505
                                                   510
Ile Asn Tyr Asn Leu Tyr Ile Leu Arg Asp Phe Lys Leu Lys Asp
```

<210> 79 <211> 528

<212> PRT <213> Saccharomyces cerevisiae

<400> 79 Ser Phe Tyr Ala Ser Pro Met Tyr Asp Phe Leu Tyr Pro Phe Arg Pro 1 10 Val Gly Asn Gln Trp Leu Pro Glu Tyr Ile Ile Phe Val Cys Ala Val 25 Ile Leu Arg Cys Thr Ile Gly Leu Gly Pro Tyr Ser Gly Lys Gly Ser Pro Pro Leu Tyr Gly Asp Phe Glu Ala Gln Arg His Trp Met Glu Ile Thr Gln His Leu Pro Leu Ser Lys Trp Tyr Trp Tyr Asp Leu Gln Tyr Trp Gly Leu Asp Tyr Pro Pro Leu Thr Ala Phe His Ser Tyr Leu Leu Gly Leu Ile Gly Ser Phe Phe Asn Pro Ser Trp Phe Ala Leu Glu Lys 100 105 Ser Arg Gly Phe Glu Ser Pro Asp Asn Gly Leu Lys Thr Tyr Met Arg 120 115 125 Ser Thr Val Ile Ile Ser Asp Ile Leu Phe Tyr Phe Pro Ala Val Ile 135 140 Tyr Phe Thr Lys Trp Leu Gly Arg Tyr Arg Asn Gln Ser Pro Ile Gly 150 155 Gln Ser Ile Ala Ala Ser Ala Ile Leu Phe Gln Pro Ser Leu Met Leu 165 170 Ile Asp His Gly His Phe Gln Tyr Asn Ser Val Met Leu Gly Leu Thr 180 185 Ala Tyr Ala Ile Asn Asn Leu Leu Asp Glu Tyr Tyr Ala Met Ala Ala 200 Val Cys Phe Val Leu Ser Ile Cys Phe Lys Gln Met Ala Leu Tyr Tyr 215 Ala Pro Ile Phe Phe Ala Tyr Leu Leu Ser Arg Ser Leu Leu Phe Pro 230 235 Lys Phe Asn Ile Ala Arg Leu Thr Val Ile Ala Phe Ala Thr Leu Ala 245 250 Thr Phe Ala Ile Ile Phe Ala Pro Leu Tyr Phe Leu Gly Gly Leu 260 265 Lys Asn Ile His Gln Cys Ile His Arg Ile Phe Pro Phe Ala Arg Gly 280 285 Ile Phe Glu Asp Lys Val Ala Asn Phe Trp Cys Val Thr Asn Val Phe 295 300 Val Lys Tyr Lys Glu Arg Phe Thr Ile Gln Gln Leu Gln Leu Tyr Ser 310 315 Leu Ile Ala Thr Val Ile Gly Phe Leu Pro Ala Met Ile Met Thr Leu 325 330 Leu His Pro Lys Lys His Leu Leu Pro Tyr Val Leu Ile Ala Cys Ser 345 Met Ser Phe Phe Leu Phe Ser Phe Gln Val His Glu Lys Thr Ile Leu 360 Ile Pro Leu Leu Pro Ile Thr Leu Leu Tyr Ser Ser Thr Asp Trp Asn 375 Val Leu Ser Leu Val Ser Trp Ile Asn Asn Val Ala Leu Phe Thr Leu 390 395 Trp Pro Leu Leu Lys Lys Asp Gly Leu His Leu Gln Tyr Ala Val Ser 410 405 Phe Leu Leu Ser Asn Trp Leu Ile Gly Asn Phe Ser Phe Ile Thr Pro 420 425

```
Asn Ser Asp Tyr Arg Arg Ser Leu Leu Pro Tyr Asn Val Val Trp
                       455
                                          460
Lys Ser Phe Ile Ile Gly Thr Tyr Ile Ala Met Gly Phe Tyr His Phe
                   470
                                      475
Leu Asp Gln Phe Val Ala Pro Pro Ser Lys Tyr Pro Asp Leu Trp Val
               485
                                  490
                                                      495
Leu Leu Asn Cys Ala Val Gly Phe Ile Cys Phe Ser Ile Phe Trp Leu
           500
                               505
Trp Ser Tyr Tyr Lys Ile Phe Thr Ser Gly Ser Lys Ser Met Lys Asp
       515
                           520
<210> 80
<211> 511
<212> PRT
<213> Pichia pastoris
<220>
<221> MOD RES
<222> (22) ... (36)
<223> Xaa is a variable amino acid
<220>
<221> MOD RES
<222> (365)...(377)
<223> Xaa is a variable amino acid
Phe Glu Asn Ser Pro Val Phe Asp Phe Leu Ser Pro Phe Gly Pro Ala
                                  10
25
Xaa Xaa Xaa Val Gly Leu Gly Ser Tyr Ser Gly Phe Asn Thr Pro
                           40
Pro Met Tyr Gly Asp Phe Glu Ala Gln Arg His Trp Met Glu Ile Thr
                      55
Gln His Leu Ser Ile Glu Lys Trp Tyr Phe Tyr Asp Leu Gln Tyr Trp
                                      7.5
                  70
Gly Leu Asp Tyr Pro Pro Leu Thr Ala Phe His Ser Tyr Phe Phe Gly
               8.5
                                  90
Lys Leu Gly Ser Phe Ile Asn Pro Ala Trp Phe Ala Leu Asp Val Ser
           100
                              105
Arg Gly Phe Glu Ser Val Asp Leu Lys Ser Tyr Met Arg Ala Thr Ala
                          120
Ile Leu Ser Glu Leu Leu Cys Phe Ile Pro Ala Val Ile Trp Tyr Cys
                      135
                                          140
Arg Trp Met Gly Leu Asn Tyr Phe Asn Gln Asn Ala Ile Glu Gln Thr
                  150
                                      155
Ile Ile Ala Ser Ala Ile Leu Phe Asn Pro Ser Leu Ile Ile Ile Asp
                                  170
               165
His Gly His Phe Gln Tyr Asn Ser Val Met Leu Gly Phe Ala Leu Leu
                              185
Ser Ile Leu Asn Leu Leu Tyr Asp Asn Phe Ala Leu Ala Ala Ile Phe
                          200
                                             205
Phe Val Leu Ser Ile Ser Phe Lys Gln Met Ala Leu Tyr Tyr Ser Pro
   210
                      215
```

Arg Phe Leu Pro Lys Ser Leu Thr Pro Gly Pro Ser Ile Ser Ser Ile 435 440 445

```
Ile Met Phe Phe Tyr Met Leu Ser Val Ser Cys Trp Pro Leu Lys Asn
                   230
                                        235
Phe Asn Leu Leu Arg Leu Ala Thr Ile Ser Ile Ala Val Leu Leu Thr
                245
                                    250
Phe Ala Thr Leu Leu Pro Phe Val Leu Val Asp Gly Met Ser Gln
                                265
Ile Gly Gln Ile Leu Phe Arg Val Phe Pro Phe Ser Arg Gly Leu Phe
                            280
                                                285
Glu Asp Lys Val Ala Asn Phe Trp Cys Thr Thr Asn Ile Leu Val Lys
                        295
                                            300
Tyr Lys Gln Leu Phe Thr Asp Lys Thr Leu Thr Arg Ile Ser Leu Val
305
                    310
                                        315
Ala Thr Leu Ile Ala Ile Ser Pro Ser Cys Phe Ile Ile Phe Thr His
                325
                                    330
Pro Lys Lys Val Leu Leu Pro Trp Ala Phe Ala Ala Cys Ser Trp Ala
                                345
Phe Tyr Leu Phe Ser Phe Gln Val His Glu Lys Ser Xaa Xaa Xaa Xaa
                            360
Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Glu Lys Asp Leu Asp Ile Ile
                        375
                                            380
Ser Met Val Cys Trp Ile Ser Asn Ile Ala Phe Phe Ser Met Trp Pro
                   390
                                        395
Leu Leu Lys Arg Asp Gly Leu Ala Leu Glu Tyr Phe Val Leu Gly Ile
               405
                                    410
Leu Ser Asn Trp Leu Ile Gly Asn Leu Asn Trp Ile Ser Lys Trp Leu
           420
                               425
                                                   430
Val Pro Ser Phe Leu Ile Pro Gly Pro Thr Leu Ser Lys Lys Val Pro
       435
                           440
                                               445
Lys Arg Asp Thr Lys Thr Val Val His Thr His Trp Phe Trp Gly Ser
                       455
                                            460
Val Thr Phe Val Ser Tyr Leu Gly Ala Thr Val Ile Gln Phe Val Asp
                   470
                                        475
Trp Leu Tyr Leu Pro Pro Ala Lys Tyr Pro Asp Leu Trp Val Ile Leu
                                   490
Asn Thr Thr Leu Ser Phe Ala Cys Phe Gly Leu Phe Trp Leu Trp
```

```
<210> 81
```

<400> 81

Phe Glu Asn Gly Ala Pro Val Gln Gln Phe Val Ser Arg Phe Arg Ser 10 Tyr Ser Ser Lys Phe Leu Phe Phe Pro Cys Leu Ile Met Ser Leu Val 20 25 Phe Met Gln Trp Leu Ile Ser Ile Gly Pro Tyr Ser Gly Tyr Asn Thr 40 Pro Pro Met Tyr Gly Asp Phe Glu Ala Gln Arg His Trp Met Glu Leu Thr Leu His Thr Pro Val Ser Gln Trp Tyr Phe Arg Asp Leu Gln Trp 70 75 Trp Gly Leu Asp Tyr Pro Pro Leu Thr Ala Tyr Val Ser Trp Phe Phe 85 90 Gly Ile Ile Gly His Tyr Phe Phe Asn Pro Glu Trp Phe Ala Asp Val 105

<211> 480

<212> PRT

<213> Schizosaccharomyces pombe

```
Thr Ser Arg Gly Phe Glu Ser Leu Glu Leu Lys Leu Phe Met Arg Ser
       115
                           120
Thr Val Ile Ala Ser His Leu Leu Ile Leu Val Pro Pro Leu Met Phe
                        135
                                           140
Tyr Ser Lys Trp Trp Ser Arg Arg Ile Pro Asn Phe Val Asp Arg Asn
                   150
                                       155
Ala Ser Leu Ile Met Val Leu Phe Gln Pro Ala Leu Leu Leu Ile Asp
               165
                                   170
His Gly His Phe Gln Tyr Asn Cys Val Met Leu Gly Leu Val Met Tyr
                               185
Ala Ile Ala Asn Leu Leu Lys Asn Gln Tyr Val Ala Ala Thr Phe Phe
                           200
Phe Cys Leu Ala Leu Thr Phe Lys Gln Met Ala Leu Tyr Phe Ala Pro
                       215
Pro Ile Phe Phe Tyr Leu Leu Gly Thr Cys Val Lys Pro Lys Ile Arg
                   230
                                       235
Phe Ser Arg Phe Ile Leu Leu Ser Val Thr Val Val Phe Thr Phe Ser
               245
                                   250
Leu Ile Leu Phe Pro Trp Ile Tyr Met Asp Tyr Lys Thr Leu Leu Pro
                               265
           260
Gln Ile Leu His Arg Val Phe Pro Phe Ala Arg Gly Leu Trp Glu Asp
       275
                           280
                                               285
Lys Val Ala Asn Phe Trp Cys Thr Leu Asn Thr Val Phe Lys Ile Arg
                       295
                                           300
Glu Val Phe Thr Leu His Gln Leu Gln Val Ile Ser Leu Ile Phe Thr
                   310
                                       315
Leu Ile Ser Ile Leu Pro Ser Cys Val Ile Leu Phe Leu Tyr Pro Arg
              325
                                  330
Lys Arg Leu Leu Ala Leu Gly Phe Ala Ser Ala Ser Trp Gly Phe Phe
          340
                               345
Leu Phe Ser Phe Gln Val His Glu Lys Ser Val Leu Leu Pro Leu Leu
                           360
Pro Thr Ser Ile Leu Cys His Gly Asn Ile Thr Thr Lys Pro Trp
                       375
                                           380
Ile Ala Leu Ala Asn Asn Leu Ala Val Phe Ser Leu Trp Pro Leu Leu
                   390
                                       395
Lys Lys Asp Gly Leu Gly Leu Gln Tyr Phe Thr Leu Val Leu Met Trp
               405
                                   410
Asn Trp Ile Gly Asp Met Val Val Phe Ser Lys Asn Val Leu Phe Arg
           420
                               425
Phe Ile Gln Leu Ser Phe Tyr Val Gly Met Ile Val Ile Leu Gly Ile
                           440
                                               445
Asp Leu Phe Ile Pro Pro Pro Ser Arg Tyr Pro Asp Leu Trp Val Ile
                    455
                                          460
Leu Asn Val Thr Leu Ser Phe Ala Gly Phe Phe Thr Ile Tyr Leu Trp
                  470
                                       475
```

```
<210> 82
```

<211> 477

<212> PRT

<213> Pichia pastoris

<220>

<221> MOD RES

<222> (329)...(341)

<223> Xaa is a variable amino acid

<400> 82 Val Gly Leu Gly Ser Tyr Ser Gly Phe Asn Thr Pro Pro Met Tyr Gly 10 Asp Phe Glu Ala Gln Arg His Trp Met Glu Ile Thr Gln His Leu Ser 25 Ile Glu Lys Trp Tyr Phe Tyr Asp Leu Gln Tyr Trp Gly Leu Asp Tyr 40 Pro Pro Leu Thr Ala Phe His Ser Tyr Phe Phe Gly Lys Leu Gly Ser 55 Phe Ile Asn Pro Ala Trp Phe Ala Leu Asp Val Ser Arg Gly Phe Glu 70 Ser Val Asp Leu Lys Ser Tyr Met Arg Ala Thr Ala Ile Leu Ser Glu 85 Leu Leu Cys Phe Ile Pro Ala Val Ile Trp Tyr Cys Arg Trp Met Gly 105 Leu Asn Tyr Phe Asn Gln Asn Ala Ile Glu Gln Thr Ile Ile Ala Ser 120 125 Ala Ile Leu Phe Asn Pro Ser Leu Ile Ile Ile Asp His Gly His Phe 135 Gln Tyr Asn Ser Val Met Leu Gly Phe Ala Leu Leu Ser Ile Leu Asn 150 155 Leu Leu Tyr Asp Asn Phe Ala Leu Ala Ile Phe Phe Val Leu Ser 165 170 Ile Ser Phe Lys Gln Met Ala Leu Tyr Tyr Ser Pro Ile Met Phe Phe 180 185 Tyr Met Leu Ser Val Ser Cys Trp Pro Leu Lys Asn Phe Asn Leu Leu 200 205 Arg Leu Ala Thr Ile Ser Ile Ala Val Leu Leu Thr Phe Ala Thr Leu 215 Leu Leu Pro Phe Val Leu Val Asp Gly Met Ser Gln Ile Gly Gln Ile 230 235 Leu Phe Arg Val Phe Pro Phe Ser Arg Gly Leu Phe Glu Asp Lys Val 250 Ala Asn Phe Trp Cys Thr Thr Asn Ile Leu Val Lys Tyr Lys Gln Leu 265 270 Phe Thr Asp Lys Thr Leu Thr Arg Ile Ser Leu Val Ala Thr Leu Ile 280 285 Ala Ile Ser Pro Ser Cys Phe Ile Ile Phe Thr His Pro Lys Lys Val 295 300 Leu Leu Pro Trp Ala Phe Ala Ala Cys Ser Trp Ala Phe Tyr Leu Phe 310 315 Ser Phe Gln Val His Glu Lys Ser Xaa Xaa Xaa Xaa Xaa Xaa Xaa 325 330 Xaa Xaa Xaa Xaa Glu Lys Asp Leu Asp Ile Ile Ser Met Val Cys 340 345 Trp Ile Ser Asn Ile Ala Phe Phe Ser Met Trp Pro Leu Leu Lys Arg 360 Asp Gly Leu Ala Leu Glu Tyr Phe Val Leu Gly Ile Leu Ser Asn Trp 375 380 Leu Ile Gly Asn Leu Asn Trp Ile Ser Lys Trp Leu Val Pro Ser Phe 390 395 Leu Ile Pro Gly Pro Thr Leu Ser Lys Lys Val Pro Lys Arg Asp Thr 405 410 Lys Thr Val Val His Thr His Trp Phe Trp Gly Ser Val Thr Phe Val 420 425 430 Ser Tyr Leu Gly Ala Thr Val Ile Gln Phe Val Asp Trp Leu Tyr Leu 440 445 Pro Pro Ala Lys Tyr Pro Asp Leu Trp Val Ile Leu Asn Thr Thr Leu

```
455
Ser Phe Ala Cys Phe Gly Leu Phe Trp Leu Trp Ile Asn
                    470
<210> 83
<211> 448
<212> PRT
<213> Drosophila melanogaster
<400> 83
Ile Ser Leu Tyr Ser Tyr Ser Gly Phe Asp Ser Pro Pro Met His Gly
Asp Tyr Glu Ala Gln Arg His Trp Gln Glu Ile Thr Val Asn Leu Ala
Val Gly Glu Trp Tyr Thr Asn Ser Ser Asn Asn Asp Leu Gln Tyr Trp
Gly Leu Asp Tyr Pro Pro Leu Thr Ala Tyr His Ser Tyr Leu Val Gly
                        55
Arg Ile Gly Ala Ser Ile Asp Pro Arg Phe Val Glu Leu His Lys Ser
                                        75
Arg Gly Phe Glu Ser Lys Glu His Lys Arg Phe Met Arg Ala Thr Val
                85
                                    90
Val Ser Ala Asp Val Leu Ile Tyr Leu Pro Ala Met Leu Leu Ala
                               105
                                                    110
Tyr Ser Leu Asp Lys Ala Phe Arg Ser Asp Asp Lys Leu Phe Leu Phe
                           120
                                               125
Thr Leu Val Ala Ala Tyr Pro Gly Gln Thr Leu Ile Asp Asn Gly His
                       135
                                           140
Phe Gln Tyr Asn Asn Ile Ser Leu Gly Phe Ala Ala Val Ala Ile Ala
                   150
                                       155
Ala Ile Leu Arg Arg Phe Tyr Ala Ala Phe Phe Phe Thr Leu
                                   170
Ala Leu Asn Tyr Lys Gln Met Glu Leu Tyr His Ser Leu Pro Phe Phe
                               185
Ala Phe Leu Leu Gly Glu Cys Val Ser Gln Lys Ser Phe Ala Ser Phe
                           200
                                                205
Ile Ala Glu Ile Ser Arg Ile Ala Ala Val Val Leu Gly Thr Phe Ala
                       215
                                            220
Ile Leu Trp Val Pro Trp Leu Gly Ser Leu Gln Ala Val Leu Gln Val
                   23.0
                                        235
Leu His Arg Leu Phe Pro Val Ala Arg Gly Val Phe Glu Asp Lys Val
               245
                                   250
Ala Asn Val Trp Cys Ala Val Asn Val Val Trp Lys Leu Lys Lys His
           260
                                265
                                                    270
Ile Ser Asn Asp Gln Met Ala Leu Val Cys Ile Ala Cys Thr Leu Ile
                           280
                                                285
Ala Ser Leu Pro Thr Asn Val Leu Leu Phe Arg Arg Arg Thr Asn Val
                       295
                                            300
Gly Phe Leu Leu Ala Leu Phe Asn Thr Ser Leu Ala Phe Phe Leu Phe
                   310
                                        315
Ser Phe Gln Val His Glu Lys Thr Ile Leu Leu Thr Ala Leu Pro Ala
               325
                                   330
Leu Phe Leu Leu Lys Cys Trp Pro Asp Glu Met Ile Leu Phe Leu Glu
                               345
                                                    350
Val Thr Val Phe Ser Met Leu Pro Leu Leu Ala Arg Asp Glu Leu Leu
                           360
Val Pro Ala Val Val Ala Thr Val Ala Phe His Leu Ile Phe Lys Cys
```

```
370
                        375
Phe Asp Ser Lys Ser Lys Leu Ser Asn Glu Tyr Pro Leu Lys Tyr Ile
                   390
                                        395
Ala Asn Ile Ser Gln Ile Leu Met Ile Ser Val Val Ala Ser Leu
               405
                                    410
Thr Val Pro Ala Pro Thr Lys Tyr Pro Asp Leu Trp Pro Leu Ile Ile
                                425
Ser Val Thr Ser Cys Gly His Phe Phe Leu Phe Phe Leu Trp Gly Asn
<210> 84
<211> 478
<212> PRT
<213> Pichia pastoris
<220>
<221> MOD RES
<222> (324)...(336)
<223> Xaa is a variable amino acid
Tyr Ser Gly Phe Asn Thr Pro Pro Met Tyr Gly Asp Phe Glu Ala Gln
                                    10
Arg His Trp Met Glu Ile Thr Gln His Leu Ser Ile Glu Lys Trp Tyr
                               25
Phe Tyr Asp Leu Gln Tyr Trp Gly Leu Asp Tyr Pro Pro Leu Thr Ala
                            40
                                                45
Phe His Ser Tyr Phe Phe Gly Lys Leu Gly Ser Phe Ile Asn Pro Ala
                        55
Trp Phe Ala Leu Asp Val Ser Arg Gly Phe Glu Ser Val Asp Leu Lys
                   70
                                        75
Ser Tyr Met Arg Ala Thr Ala Ile Leu Ser Glu Leu Leu Cys Phe Ile
                                    90
Pro Ala Val Ile Trp Tyr Cys Arg Trp Met Gly Leu Asn Tyr Phe Asn
                               105
Gln Asn Ala Ile Glu Gln Thr Ile Ile Ala Ser Ala Ile Leu Phe Asn
                           120
                                                125
Pro Ser Leu Ile Ile Asp His Gly His Phe Gln Tyr Asn Ser Val
                       135
                                            140
Met Leu Gly Phe Ala Leu Leu Ser Ile Leu Asn Leu Leu Tyr Asp Asn
                   150
                                        155
Phe Ala Leu Ala Ala Ile Phe Phe Val Leu Ser Ile Ser Phe Lys Gln
               165
                                    170
Met Ala Leu Tyr Tyr Ser Pro Ile Met Phe Phe Tyr Met Leu Ser Val
           180
                               185
                                                    190
Ser Cys Trp Pro Leu Lys Asn Phe Asn Leu Leu Arg Leu Ala Thr Ile
       195
                           200
                                                205
Ser Ile Ala Val Leu Leu Thr Phe Ala Thr Leu Leu Leu Pro Phe Val
                       215
Leu Val Asp Gly Met Ser Gln Ile Gly Gln Ile Leu Phe Arg Val Phe
                   230
                                        235
Pro Phe Ser Arg Gly Leu Phe Glu Asp Lys Val Ala Asn Phe Trp Cys
                                   250
Thr Thr Asn Ile Leu Val Lys Tyr Lys Gln Leu Phe Thr Asp Lys Thr
           260
                                                270
                               265
Leu Thr Arg Ile Ser Leu Val Ala Thr Leu Ile Ala Ile Ser Pro Ser
```

```
Cys Phe Ile Ile Phe Thr His Pro Lys Lys Val Leu Leu Pro Trp Ala
                      295
Phe Ala Ala Cys Ser Trp Ala Phe Tyr Leu Phe Ser Phe Gln Val His
                  310
                                     315
325
                                 330
Glu Lys Asp Leu Asp Ile Ile Ser Met Val Cys Trp Ile Ser Asn Ile
                             345
Ala Phe Phe Ser Met Trp Pro Leu Leu Lys Arg Asp Gly Leu Ala Leu
       355
                          360
Glu Tyr Phe Val Leu Gly Ile Leu Ser Asn Trp Leu Ile Gly Asn Leu
   370
                      375
Asn Trp Ile Ser Lys Trp Leu Val Pro Ser Phe Leu Ile Pro Gly Pro
                  390
Thr Leu Ser Lys Lys Val Pro Lys Arg Asp Thr Lys Thr Val Val His
                                 410
Thr His Trp Phe Trp Gly Ser Val Thr Phe Val Ser Tyr Leu Gly Ala
                             425
Thr Val Ile Gln Phe Val Asp Trp Leu Tyr Leu Pro Pro Ala Lys Tyr
                          440
                                            445
Pro Asp Leu Trp Val Ile Leu Asn Thr Thr Leu Ser Phe Ala Cys Phe
                     455
Gly Leu Phe Trp Leu Trp Ile Asn Tyr Asn Leu Tyr Ile Leu
                  470
```

<210> 85

<211> 459

<212> PRT

<213> Arabidopsis thaliana

<400> 85

Tyr Ser Gly Ala Gly Ile Pro Pro Lys Phe Gly Asp Phe Glu Ala Gln Arg His Trp Met Glu Ile Thr Thr Asn Leu Pro Val Ile Asp Trp Tyr Arg Asn Gly Thr Tyr Asn Asp Leu Thr Tyr Trp Gly Leu Asp Tyr Pro Pro Leu Thr Ala Tyr Gln Ser Tyr Ile His Gly Ile Phe Leu Arg Phe 55 60 Phe Asn Pro Glu Ser Val Ala Leu Leu Ser Ser Arg Gly His Glu Ser 70 75 Tyr Leu Gly Lys Leu Leu Met Arg Trp Thr Val Leu Ser Ser Asp Ala 90 Phe Ile Phe Pro Ala Ala Leu Phe Phe Val Leu Val Tyr His Arg 100 105 110 Asn Arg Thr Arg Gly Gly Lys Ser Glu Val Ala Trp His Ile Ala Met 120 125 Ile Leu Leu Asn Pro Cys Leu Ile Leu Ile Asp His Gly His Phe Gln 135 Tyr Asn Cys Ile Ser Leu Gly Leu Thr Val Gly Ala Ile Ala Ala Val 150 155 Leu Cys Glu Ser Glu Val Leu Thr Cys Val Leu Phe Ser Leu Ala Leu 170 Ser His Lys Gln Met Ser Ala Tyr Phe Ala Pro Ala Phe Phe Ser His 185 190 Leu Leu Gly Lys Cys Leu Arg Arg Lys Ser Pro Ile Leu Ser Val Ile

<211> 277 <212> PRT

```
Lys Leu Gly Ile Ala Val Ile Val Thr Phe Val Ile Phe Trp Trp Pro
                        215
Tyr Val His Ser Leu Asp Asp Phe Leu Met Val Leu Ser Arg Leu Ala
                    230
                                         235
Pro Phe Glu Arg Gly Ile Tyr Glu Asp Tyr Val Ala Asn Phe Trp Cys
                245
                                     250
                                                         255
Thr Thr Ser Ile Leu Ile Lys Trp Lys Asn Leu Phe Thr Thr Gln Ser
                                 265
Leu Lys Ser Ile Ser Leu Ala Ala Thr Ile Leu Ala Ser Leu Pro Ser
                             280
Met Val Gln Gln Ile Leu Ser Pro Ser Asn Glu Gly Phe Leu Tyr Gly
    290
                        295
                                             300
Leu Leu Asn Ser Ser Met Ala Phe Tyr Leu Phe Ser Phe Gln Val His
305
                    310
                                         315
Glu Lys Ser Ile Leu Met Pro Phe Leu Ser Ala Thr Leu Leu Ala Leu
                                     330
Lys Leu Pro Asp His Phe Ser His Leu Thr Tyr Tyr Ala Leu Phe Ser
            340
                                345
                                                     350
Met Phe Pro Leu Leu Cys Arg Asp Lys Leu Leu Ile Pro Tyr Leu Thr
                            360
                                                 365
Leu Ser Phe Leu Phe Thr Val Ile Tyr His Ser Pro Gly Asn His His
                        375
                                             380
Ala Ile Gln Lys Thr Asp Val Ser Phe Phe Ser Phe Lys Asn Phe Pro
                    390
                                         395
Gly Tyr Val Phe Leu Leu Arg Thr His Phe Phe Ile Ser Val Val Leu
                405
                                     410
                                                         415
His Val Leu Tyr Leu Thr Ile Lys Pro Pro Gln Lys Tyr Pro Phe Leu
            420
                                425
Phe Glu Ala Leu Ile Met Ile Leu Cys Phe Ser Tyr Phe Ile Met Phe
                            440
Ala Phe Tyr Thr Asn Tyr Thr Gln Trp Thr Leu
                        455
<210> 86
<211> 836
<212> DNA
<213> Kluyveromyces lactis
<400> 86
atctctgttt caacagctct tgcattcatt ggttctttcg gtccaatcta tatctttgga 60
ggatacaaga acttagtgca atcaatgcac aggatttttc catttgccag gggtatcttt 120
gaagataaag ttgcgaattt ttggtgcgtt tctaatattt tcatcaaata tagaaatcta 180
ttcactcaga aggatettea attataetea ttaetegeaa eagttattgg gettttaeea 240
tcattcatta taacattttt atacccgaag agacatttac taccatatgc tttggccgca 300
tgttcgatgt cattettett atteagette caggtteatg aaaagacaat ettattaeet 360
ttacttccta ttacactctt gtacacgtca agagattgga atgttctatc attggtttgt 420
tggattaaca acgtggcatt gtttacactc tggccattac tgaaaaagga caatctagta 480
ttgcaatatg gagtcatgtt catgtttagc aattggttga tcggtaactt cagtttcgtc 540
acaccacgct tcctcccaaa atttttgaca ccagggccat ccatcagtga tatagatgtt 600
gattatagac gggcaagttt actacccaag agcctaatat ggagattaat cattgttggc 660
teatatattg caatggggat tatteatttt etagaetatt aegteteece gecateaaaa 720
taccctgatt tatgggtgct tgccaattgt tccttgggct tctcatgttt tgtgacattt 780
tggatatgga acaattataa ttattcgaaa tgagaaacag cactttgcaa gattta
<210> 87
```

```
<213> Kluyveromyces lactis
<400> 87
Ile Ser Val Ser Thr Ala Leu Ala Phe Ile Gly Ser Phe Gly Pro Ile
                                    10
Tyr Ile Phe Gly Gly Tyr Lys Asn Leu Val Gln Ser Met His Arg Ile
                                25
Phe Pro Phe Ala Arg Gly Ile Phe Glu Asp Lys Val Ala Asn Phe Trp
Cys Val Ser Asn Ile Phe Ile Lys Tyr Arg Asn Leu Phe Thr Gln Lys
                        55
Asp Leu Gln Leu Tyr Ser Leu Leu Ala Thr Val Ile Gly Leu Leu Pro
                    70
Ser Phe Ile Ile Thr Phe Leu Tyr Pro Lys Arg His Leu Leu Pro Tyr
                                    90
Ala Leu Ala Ala Cys Ser Met Ser Phe Phe Leu Phe Ser Phe Gln Val
                                105
His Glu Lys Thr Ile Leu Leu Pro Leu Pro Ile Thr Leu Leu Tyr
                            120
                                                 125
Thr Ser Arg Asp Trp Asn Val Leu Ser Leu Val Cys Trp Ile Asn Asn
                        135
                                            140
Val Ala Leu Phe Thr Leu Trp Pro Leu Leu Lys Lys Asp Asn Leu Val
                    150
                                        155
Leu Gln Tyr Gly Val Met Phe Met Phe Ser Asn Trp Leu Ile Gly Asn
                165
                                   170
                                                         175
Phe Ser Phe Val Thr Pro Arg Phe Leu Pro Lys Phe Leu Thr Pro Gly
            180
                                185
Pro Ser Ile Ser Asp Ile Asp Val Asp Tyr Arg Arg Ala Ser Leu Leu
                           200
                                                205
Pro Lys Ser Leu Ile Trp Arg Leu Ile Ile Val Gly Ser Tyr Ile Ala
                        215
Met Gly Ile Ile His Phe Leu Asp Tyr Tyr Val Ser Pro Pro Ser Lys
                   230
                                        235
Tyr Pro Asp Leu Trp Val Leu Ala Asn Cys Ser Leu Gly Phe Ser Cys
                                   250
Phe Val Thr Phe Trp Ile Trp Asn Asn Tyr Asn Tyr Ser Lys Glu Thr
                                265
Ala Leu Cys Lys Ile
       275
<210> 88
<211> 284
<212> PRT
<213> Kluyveromyces lactis
<220>
<221> MOD RES
<222> (116)...(127)
<223> Xaa is a variable amino acid
<220>
<221> MOD_RES
<222> 271
<223> Xaa is a variable amino acid
Ile Ser Val Ser Thr Ala Leu Ala Phe Ile Gly Ser Phe Gly Pro Ile
```

```
Tyr Ile Phe Gly Gly Tyr Lys Asn Leu Val Gln Ser Met His Arg Ile
                              25
Phe Pro Phe Ala Arg Gly Ile Phe Glu Asp Lys Val Ala Asn Phe Trp
Cys Val Ser Asn Ile Phe Ile Lys Tyr Arg Asn Leu Phe Thr Gln Lys
                      55
Asp Leu Gln Leu Tyr Ser Leu Leu Ala Thr Val Ile Gly Leu Leu Pro
                   70
Ser Phe Ile Ile Thr Phe Leu Tyr Pro Lys Arg His Leu Leu Pro Tyr
                                  90
Ala Leu Ala Ala Cys Ser Met Ser Phe Phe Leu Phe Ser Phe Gln Val
120
Thr Ser Arg Asp Trp Asn Val Leu Ser Leu Val Cys Trp Ile Asn Asn
                      135
                                          140
Val Ala Leu Phe Thr Leu Trp Pro Leu Leu Lys Lys Asp Asn Leu Val
                  150
                                      155
Leu Gln Tyr Gly Val Met Phe Met Phe Ser Asn Trp Leu Ile Gly Asn
               165
                                  170
Phe Ser Phe Val Thr Pro Arg Phe Leu Pro Lys Phe Leu Thr Pro Gly
           180
                              185
Pro Ser Ile Ser Asp Ile Asp Val Asp Tyr Arg Arg Ala Ser Leu Leu
                          200
                                             205
Pro Lys Ser Leu Ile Trp Arg Leu Ile Ile Val Gly Ser Tyr Ile Ala
                      215
                                          220
Met Gly Ile Ile His Phe Leu Asp Tyr Tyr Val Ser Pro Pro Ser Gln
                  230
                                     235
Glu Arg Tyr Lys Tyr Pro Asp Leu Trp Val Leu Ala Asn Cys Ser Leu
                                 250
Gly Phe Ser Cys Phe Val Thr Phe Trp Ile Trp Asn Asn Tyr Xaa Leu
                             265
Phe Glu Arg Met Arg Asn Ser Thr Leu Gln Asp Leu
```

<210> 89

<211> 280

<212> PRT

<213> Saccharomyces cerevisiae

<400> 89

 Ile Ala Phe Ala Phe Ala Thr Leu 5
 Ala Thr Phe Ala Thr Phe Ala Ile Ile Phe Ala Pro Leu 10
 Ala Phe Ala Pro Leu 15
 Ala Ile Phe Ala Ile Phe Ala Pro Ile Phe Ala Arg Arg 25
 Asn Ile His Gln Cys Ile His Arg 30
 Arg 30

 Ile Phe Pro Phe Ala Arg Gly Ile Phe Glu Asp Lys Val Ala Asn Phe 35
 Ala Phe Val Lys Tyr Lys Glu Arg Phe Thr Ile 50
 Ala Phe Val Lys Tyr Lys Glu Arg Phe Thr Ile Eu Eu Fro S5
 Ala Phe Phe Phe Leu Phe Eu Fro S6

 Gln Gln Leu Gln Leu Tyr Ser Leu Ile Ala Thr Val Ile Gly Phe Leu S5
 Ala Phe Phe Phe Leu Phe Phe Leu Phe Ser Phe Gln S5

 Pro Ala Met Ile Met Thr Leu Leu Leu His Pro Lys Lys His Leu Leu Pro S5

 Tyr Val Leu Ile Ala Cys Ser Met Ser Phe Phe Leu Phe Ser Phe Gln 100

 Val His Glu Lys Thr Ile Leu Ile Pro Leu Leu Pro Ile Thr Leu Leu

```
120
Tyr Ser Ser Thr Asp Trp Asn Val Leu Ser Leu Val Ser Trp Ile Asn
                    135
                                           140
Asn Val Ala Leu Phe Thr Leu Trp Pro Leu Leu Lys Lys Asp Gly Leu
                   150
                                       155
His Leu Gln Tyr Ala Val Ser Phe Leu Leu Ser Asn Trp Leu Ile Gly
                                  170
               165
Asn Phe Ser Phe Ile Thr Pro Arg Phe Leu Pro Lys Ser Leu Thr Pro
            180
                               185
Gly Pro Ser Ile Ser Ser Ile Asn Ser Asp Tyr Arg Arg Ser Leu
                           200
Leu Pro Tyr Asn Val Val Trp Lys Ser Phe Ile Ile Gly Thr Tyr Ile
                       215
Ala Met Gly Phe Tyr His Phe Leu Asp Gln Phe Val Ala Pro Pro Ser
                   230
                                       235
Lys Tyr Pro Asp Leu Trp Val Leu Leu Asn Cys Ala Val Gly Phe Ile
               245
                                   250
Cys Phe Ser Ile Phe Trp Leu Trp Ser Tyr Tyr Lys Ile Phe Thr Ser
                               265
Gly Ser Lys Ser Met Lys Asp Leu
<210> 90
<211> 284
<212> PRT
<213> Kluyveromyces lactis
<220>
<221> MOD RES
<222> (116)...(127)
<223> Xaa is a variable amino acid
<220>
<221> MOD RES
<222> 271
<223> Xaa is a variable amino acid
<400> 90
Ile Ser Val Ser Thr Ala Leu Ala Phe Ile Gly Ser Phe Gly Pro Ile
                                  10
Tyr Ile Phe Gly Gly Tyr Lys Asn Leu Val Gln Ser Met His Arg Ile
                               25
Phe Pro Phe Ala Arg Gly Ile Phe Glu Asp Lys Val Ala Asn Phe Trp
                           40
                                               45
Cys Val Ser Asn Ile Phe Ile Lys Tyr Arg Asn Leu Phe Thr Gln Lys
                       55
                                           60
Asp Leu Gln Leu Tyr Ser Leu Leu Ala Thr Val Ile Gly Leu Leu Pro
                   70
                                       75
Ser Phe Ile Ile Thr Phe Leu Tyr Pro Lys Arg His Leu Leu Pro Tyr
                                  90
Ala Leu Ala Ala Cys Ser Met Ser Phe Phe Leu Phe Ser Phe Gln Val
           100
                               105
120
                                              125
Thr Ser Arg Asp Trp Asn Val Leu Ser Leu Val Cys Trp Ile Asn Asn
                       135
                                          140
Val Ala Leu Phe Thr Leu Trp Pro Leu Leu Lys Lys Asp Asn Leu Val
```

```
150
Leu Gln Tyr Gly Val Met Phe Met Phe Ser Asn Trp Leu Ile Gly Asn
                                    170
Phe Ser Phe Val Thr Pro Arg Phe Leu Pro Lys Phe Leu Thr Pro Gly
                                185
Pro Ser Ile Ser Asp Ile Asp Val Asp Tyr Arg Arg Ala Ser Leu Leu
                            200
Pro Lys Ser Leu Ile Trp Arg Leu Ile Ile Val Gly Ser Tyr Ile Ala
                       215
                                            220
Met Gly Ile Ile His Phe Leu Asp Tyr Tyr Val Ser Pro Pro Ser Gln
                    230
                                        235
Glu Arg Tyr Lys Tyr Pro Asp Leu Trp Val Leu Ala Asn Cys Ser Leu
                245
                                    250
Gly Phe Ser Cys Phe Val Thr Phe Trp Ile Trp Asn Asn Tyr Xaa Leu
                                265
Phe Glu Arg Met Arg Asn Ser Thr Leu Gln Asp Leu
        275
<210> 91
<211> 250
<212> PRT
<213> Schizosaccharomyces pombe
<400> 91
Leu Ser Val Thr Val Val Phe Thr Phe Ser Leu Ile Leu Phe Pro Trp
                                   10
Ile Tyr Met Asp Tyr Lys Thr Leu Leu Pro Gln Ile Leu His Arg Val
                               25
Phe Pro Phe Ala Arg Gly Leu Trp Glu Asp Lys Val Ala Asn Phe Trp
                           40
Cys Thr Leu Asn Thr Val Phe Lys Ile Arg Glu Val Phe Thr Leu His
Gln Leu Gln Val Ile Ser Leu Ile Phe Thr Leu Ile Ser Ile Leu Pro
Ser Cys Val Ile Leu Phe Leu Tyr Pro Arg Lys Arg Leu Leu Ala Leu
Gly Phe Ala Ser Ala Ser Trp Gly Phe Phe Leu Phe Ser Phe Gln Val
                               105
His Glu Lys Ser Val Leu Leu Pro Leu Leu Pro Thr Ser Ile Leu Leu
                           120 ·
                                               125
Cys His Gly Asn Ile Thr Thr Lys Pro Trp Ile Ala Leu Ala Asn Asn
                       135
                                            140
Leu Ala Val Phe Ser Leu Trp Pro Leu Leu Lys Lys Asp Gly Leu Gly
                   150
                                       155
Leu Gln Tyr Phe Thr Leu Val Leu Met Trp Asn Trp Ile Gly Asp Met
                                   170
               165
Val Val Phe Ser Lys Asn Val Leu Phe Arg Phe Ile Gln Leu Ser Phe
           180
                                185
Tyr Val Gly Met Ile Val Ile Leu Gly Ile Asp Leu Phe Ile Pro Pro
       195
                            200
Pro Ser Arg Tyr Pro Asp Leu Trp Val Ile Leu Asn Val Thr Leu Ser
                       215
                                            220
Phe Ala Gly Phe Phe Thr Ile Tyr Leu Trp Thr Leu Gly Arg Leu Leu
                   230
                                        235
His Ile Ser Ser Lys Leu Ser Thr Asp Leu
```

245

```
<210> 92
<211> 238
<212> PRT
<213> Kluyveromyces lactis
<220>
<221> MOD RES
<222> (88)...(99)
<223> Xaa is a variable amino acid
<400> 92
Met His Arg Ile Phe Pro Phe Ala Arg Gly Ile Phe Glu Asp Lys Val
Ala Asn Phe Trp Cys Val Ser Asn Ile Phe Ile Lys Tyr Arg Asn Leu
Phe Thr Gln Lys Asp Leu Gln Leu Tyr Ser Leu Leu Ala Thr Val Ile
                            40
Gly Leu Leu Pro Ser Phe Ile Ile Thr Phe Leu Tyr Pro Lys Arg His
                        55
Leu Leu Pro Tyr Ala Leu Ala Ala Cys Ser Met Ser Phe Phe Leu Phe
                    70
                                        75
Ser Phe Gln Val His Glu Lys Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa
                85
                                    90
Xaa Xaa Xaa Tyr Thr Ser Arg Asp Trp Asn Val Leu Ser Leu Val Cys
            100
                               105
                                                    110
Trp Ile Asn Asn Val Ala Leu Phe Thr Leu Trp Pro Leu Leu Lys Lys
                           120
                                                125
Asp Asn Leu Val Leu Gln Tyr Gly Val Met Phe Met Phe Ser Asn Trp
                       135
                                            140
Leu Ile Gly Asn Phe Ser Phe Val Thr Pro Arg Phe Leu Pro Lys Phe
                   150
                                        155
Leu Thr Pro Gly Pro Ser Ile Ser Asp Ile Asp Val Asp Tyr Arg Arg
                                    170
Ala Ser Leu Leu Pro Lys Ser Leu Ile Trp Arg Leu Ile Ile Val Gly
                                185
Ser Tyr Ile Ala Met Gly Ile Ile His Phe Leu Asp Tyr Tyr Val Ser
                            200
                                                205
Pro Pro Ser Lys Tyr Pro Asp Leu Trp Val Leu Ala Asn Cys Ser Leu
                       215
                                            220
Gly Phe Ser Cys Phe Val Thr Phe Trp Ile Trp Asn Asn Tyr
                    230
                                        235
<210> 93
<211> 219
<212> PRT
<213> Arabidopsis thaliana
<400> 93
Leu Ser Arg Leu Ala Pro Phe Glu Arg Gly Ile Tyr Glu Asp Tyr Val
                                    10
Ala Asn Phe Trp Cys Thr Thr Ser Ile Leu Ile Lys Trp Lys Asn Leu
                                25
Phe Thr Thr Gln Ser Leu Lys Ser Ile Ser Leu Ala Ala Thr Ile Leu
                            40
                                                4.5
Ala Ser Leu Pro Ser Met Val Gln Gln Ile Leu Ser Pro Ser Asn Glu
```

```
Gly Phe Leu Tyr Gly Leu Leu Asn Ser Ser Met Ala Phe Tyr Leu Phe
                                       75
Ser Phe Gln Val His Glu Lys Ser Ile Leu Met Pro Phe Leu Ser Ala
                                   90
Thr Leu Leu Ala Leu Lys Leu Pro Asp His Phe Ser His Leu Thr Tyr
                               105
Tyr Ala Leu Phe Ser Met Phe Pro Leu Leu Cys Arg Asp Lys Leu Leu
                           120
Ile Pro Tyr Leu Thr Leu Ser Phe Leu Phe Thr Val Ile Tyr His Ser
                       135
Pro Gly Asn His His Ala Ile Gln Lys Thr Asp Val Ser Phe Phe Ser
                   150
                                       155
Phe Lys Asn Phe Pro Gly Tyr Val Phe Leu Leu Arg Thr His Phe Phe
                                   170
Ile Ser Val Val Leu His Val Leu Tyr Leu Thr Ile Lys Pro Pro Gln
                               185
                                                   190
Lys Tyr Pro Phe Leu Phe Glu Ala Leu Ile Met Ile Leu Cys Phe Ser
                           200
Tyr Phe Ile Met Phe Ala Phe Tyr Thr Asn Tyr
                       215 -
<210> 94
<211> 252
<212> PRT
<213> Kluyveromyces lactis
<220>
<221> MOD RES
<222> (114)...(125)
<223> Xaa is a variable amino acid
<400> 94
Val Ser Thr Ala Leu Ala Phe Ile Gly Ser Phe Gly Pro Ile Tyr Ile
                                  10
Phe Gly Gly Tyr Lys Asn Leu Val Gln Ser Met His Arg Ile Phe Pro
                               25
Phe Ala Arg Gly Ile Phe Glu Asp Lys Val Ala Asn Phe Trp Cys Val
Ser Asn Ile Phe Ile Lys Tyr Arg Asn Leu Phe Thr Gln Lys Asp Leu
                       55
Gln Leu Tyr Ser Leu Leu Ala Thr Val Ile Gly Leu Leu Pro Ser Phe
                  70
                                      75
Ile Ile Thr Phe Leu Tyr Pro Lys Arg His Leu Leu Pro Tyr Ala Leu
               85
                                  90
Ala Ala Cys Ser Met Ser Phe Phe Leu Phe Ser Phe Gln Val His Glu
           100
                               105
120
                                              125
Arg Asp Trp Asn Val Leu Ser Leu Val Cys Trp Ile Asn Asn Val Ala
                       135
                                          140
Leu Phe Thr Leu Trp Pro Leu Leu Lys Lys Asp Asn Leu Val Leu Gln
                  150
                                      155
Tyr Gly Val Met Phe Met Val Thr Pro Arg Phe Leu Pro Lys Phe Leu
              165
                                  170
Thr Pro Gly Pro Ser Ile Ser Asp Ile Asp Val Asp Tyr Arg Arg Ala
           180
                              185
Ser Leu Leu Pro Lys Ser Leu Ile Trp Arg Leu Ile Ile Val Gly Ser
```

```
200
Tyr Ile Ala Met Gly Ile Ile His Phe Leu Asp Tyr Tyr Val Ser Pro
                      215
                                            220
Pro Ser Lys Tyr Pro Asp Leu Trp Val Leu Ala Asn Cys Ser Leu Gly
                   230
                                       235
Phe Ser Cys Phe Val Thr Phe Trp Ile Trp Asn Asn
               245
<210> 95
<211> 259
<212> PRT
<213> Homo sapiens
<400> 95
Val Lys Leu Ala Cys Ile Val Val Ala Ser Phe Val Leu Cys Trp Leu
                                   10
Pro Phe Phe Thr Glu Arg Glu Gln Thr Leu Gln Val Leu Arg Arg Leu
                                25
Phe Pro Val Asp Arg Gly Leu Phe Glu Asp Lys Val Ala Asn Ile Trp
                                               45
Cys Ser Phe Asn Val Phe Leu Lys Ile Lys Asp Ile Leu Pro Arg His
                       55
Ile Gln Leu Ile Met Ser Phe Cys Phe Thr Phe Leu Ser Leu Leu Pro
                   70
                                       75
Ala Cys Ile Lys Leu Ile Leu Gln Pro Ser Ser Lys Gly Phe Lys Phe
              85
                                   90
Thr Leu Val Ser Cys Ala Leu Ser Phe Phe Leu Phe Ser Phe Gln Val
                               105
His Glu Lys Ser Ile Leu Leu Val Ser Leu Pro Val Cys Leu Val Leu
                          120
Ser Glu Ile Pro Phe Met Ser Thr Trp Phe Leu Leu Val Ser Thr Phe
                       135
Ser Met Leu Pro Leu Leu Lys Asp Glu Leu Leu Met Pro Ser Val
                   150
                                       155
Val Thr Thr Met Ala Phe Phe Ile Ala Cys Val Thr Ser Phe Ser Ile
               165
                                   170
Phe Glu Lys Thr Ser Glu Glu Glu Leu Gln Leu Lys Ser Phe Ser Ile
                               185
                                                    190
Ser Val Arg Lys Tyr Leu Pro Cys Phe Thr Phe Leu Ser Arg Ile Ile
                           200
                                               205
Gln Tyr Leu Phe Leu Ile Ser Val Ile Thr Met Val Leu Leu Thr Leu
                       215
                                           220
Met Thr Val Thr Leu Asp Pro Pro Gln Lys Leu Pro Asp Leu Phe Ser
                   230
                                      235
Val Leu Val Cys Phe Val Ser Cys Leu Asn Phe Leu Phe Phe Leu Val
               245
                                   250
Tyr Phe Asn
<210> 96
<211> 1617
<212> DNA
<213> Mus musculus
```

<400> 96

atgaagatga gacgctacaa gctctttctc atgttctgta tggctggcct gtgcctcata 60

```
teetteetge aettetttaa gaeettatee tatgteaeet teeegagaga aetggeetee 120
ctcagcccta acctcgtatc cagcttcttc tggaacaatg cccctgtcac tccccaggcc 180
agtccggagc cgggtggccc cgacctattg cggacacccc tctactccca ctctcccctg 240
ctccagccac tgtccccgag caaggccaca gaggaactgc accgggtgga cttcgtgttg 300
ccggaggaca ccacggagta ttttgtgcgc accaaagctg gtggtgtgtg cttcaaacca 360
ggtaccagga tgctggagaa accttcgcca gggcggacag aggagaagcc cgaagtgtct 420
gagggeteet eageeegggg acetgetegg aggeeeatga ggeaegtgtt gagtaegegg 480
gagegeetgg geageegggg cactaggege aagtgggttg agtgtgtgtg cetgeeagge 540
tggcacgggc ccagttgcgg ggtgcccacg gtggtgcagt attccaacct gcccaccaag 600
gaacgeetgg tacceaggga ggtacegagg egggttatea aegeeateaa cateaaceae 660
gagttcgacc tgctggatgt gcgcttccat gagctgggag atgttgtgga cgccttcgtg 720
gtctgtgaat ctaatttcac cgcctacggg gagcctcggc cgctcaagtt ccgagagatg 780
ctgaccaatg gcaccttcga gtacatccgc cacaaggtgc tctatgtctt cctggaccat 840
ttcccacctg gtggccgtca ggacggctgg attgcggatg actacctgcg caccttcctc 900
acccaggatg gcgtctcccg cctgcgcaac ctgcggcccg atgacgtctt tatcatcgac 960
gatgcggacg agatccctgc gcgtgatggt gtgctgttcc tcaaactcta cgatggctgg 1020
acagageeet tegeetteea eatgeggaag teeetgtatg gtttettetg gaageageeg 1080
ggcacactgg aggtggtgtc aggctgcacc atggacatgc tgcaggccgt gtatgggctg 1140
gatggcatcc gcctgcgccg ccgccagtac tacaccatgc ccaacttccg gcagtatgag 1200
aaccgcaccg gccacatcct agtgcagtgg tctctcggca gcccctgca cttcgcgggc 1260
tggcattgct cctggtgctt cacacccgag ggcatctact ttaaactcgt gtcagcccag 1320
aatggcgact tcccccgctg gggtgactat gaggacaaga gggacctcaa ttacatccgc 1380
agettgatee geactggggg atggttegae ggaacgeage aggagtacee teetgeggae 1440
cccagtgagc acatgtatgc tcctaaatac ctgctcaaga actatgacca gttccgctac 1500
ttgctggaaa atccctaccg ggagcccaag agcactgtag agggtgggcg ccagaaccag 1560
ggctcagatg gaaggccatc tgctgtcagg ggcaagttgg atacagtgga gggctag
```

<210> 97 <211> 536

<212> PRT <213> Mus musculus

<400> 97

Met Arg Arg Tyr Lys Leu Phe Leu Met Phe Cys Met Ala Gly Leu Cys 10 Leu Ile Ser Phe Leu His Phe Phe Lys Thr Leu Ser Tyr Val Thr Phe 25 Pro Arg Glu Leu Ala Ser Leu Ser Pro Asn Leu Ile Ser Ser Phe Phe 40 Trp Asn Asn Ala Pro Val Thr Pro Gln Ala Ser Pro Glu Pro Gly Asp 55 Pro Asp Leu Leu Arg Thr Pro Leu Tyr Ser His Ser Pro Leu Leu Gln 70 75 Pro Leu Ser Pro Ser Lys Ala Thr Glu Glu Leu His Arg Val Asp Phe 90 Val Leu Pro Glu Asp Thr Thr Glu Tyr Phe Val Arg Thr Lys Ala Gly 105 Gly Val Cys Phe Lys Pro Gly Thr Arg Met Leu Glu Lys Pro Ser Pro 120 Gly Arg Thr Glu Glu Lys Thr Glu Val Ser Glu Gly Ser Ser Ala Arg 135 Gly Pro Ala Arg Arg Pro Met Arg His Val Leu Ser Ser Arg Glu Arg 150 155 Leu Gly Ser Arg Gly Thr Arg Arg Lys Trp Val Glu Cys Val Cys Leu 165 170 175 Pro Gly Trp His Gly Pro Ser Cys Gly Val Pro Thr Val Val Gln Tyr 185 Ser Asn Leu Pro Thr Lys Glu Arg Leu Val Pro Arg Glu Val Pro Arg

```
230
                                         235
Asp Ser Asn Phe Thr Ala Tyr Gly Glu Pro Arg Pro Leu Lys Phe Arg
                245
                                     250
Glu Met Leu Thr Asn Gly Thr Phe Glu Tyr Ile Arg His Lys Val Leu
                                 265
Tyr Val Phe Leu Asp His Phe Pro Pro Gly Gly Arg Gln Asp Gly Trp
                             280
Ile Ala Asp Asp Tyr Leu Arg Thr Phe Leu Thr Gln Asp Gly Val Ser
                         295
Arg Leu Arg Asn Leu Arg Pro Asp Asp Val Phe Ile Ile Asp Asp Ala
                    310
                                         315
Asp Glu Ile Pro Ala Arg Asp Gly Val Leu Phe Leu Lys Leu Tyr Asp
                325
                                     330
                                                          335
Gly Trp Thr Glu Pro Phe Ala Phe His Met Arg Lys Ser Leu Tyr Gly
                                 345
                                                      350
Phe Phe Trp Lys Gln Pro Gly Thr Leu Glu Val Val Ser Gly Cys Thr
        355
                             360
                                                  365
Met Asp Met Leu Gln Ala Val Tyr Gly Leu Asp Gly Ile Arg Leu Arg
                        375
                                             380
Arg Arg Gln Tyr Tyr Thr Met Pro Asn Phe Arg Gln Tyr Glu Asn Arg
                    390
                                         395
Thr Gly His Ile Leu Val Gln Trp Ser Leu Gly Ser Pro Leu His Phe
                405
                                     410
Ala Gly Trp His Cys Ser Trp Cys Phe Thr Pro Glu Gly Ile Tyr Phe
            420
                                 425
Lys Leu Val Ser Ala Gln Asn Gly Asp Phe Pro Arg Trp Gly Asp Tyr
                            440
Glu Asp Lys Arg Asp Leu Asn Tyr Ile Arg Ser Leu Ile Arg Thr Gly
                        455
                                             460
Gly Trp Phe Asp Gly Thr Gln Glu Tyr Pro Pro Ala Asp Pro Ser
                    470
                                         475
Glu His Met Tyr Ala Pro Lys Tyr Leu Leu Lys Asn Tyr Asp Gln Phe
                485
                                     490
                                                          495
Arg Tyr Leu Leu Glu Asn Pro Tyr Arg Glu Pro Lys Ser Thr Val Glu
            500
                                 505
                                                      510
Gly Gly Arg Gln Asn Gln Gly Ser Asp Gly Arg Ser Ser Ala Val Arg
                            520
                                                 525
Gly Lys Leu Asp Thr Ala Glu Gly
    530
                        535
<210> 98
<211> 2115
<212> DNA
<213> Homo sapiens
<400> 98
gaaatgaacc tetettattg attittattg geetagagee aggagtaetg catteagttg 60
actttcaggg taaaaagaaa acagtcctgg ttgttgtcat cataaacata tggaccagtg 120
tgatggtgaa atgagatgag gctccgcaat ggaactgtag ccactgcttt agcatttatc 180
actteettee ttaetttgte ttggtataet acatggcaaa atgggaaaga aaaactgatt 240
gcttatcaac gagaattcct tgctttgaaa gaacgtcttc gaatagctga acacagaatc 300
```

195 200 205 Arg Val Ile Asn Ala Ile Asn Ile Asn His Glu Phe Asp Leu Leu Asp

Val Arg Phe His Glu Leu Gly Asp Val Val Asp Ala Phe Val Val Cys

220

215

tcacageget ettetgaatt aaataegatt gtgeaacagt teaagegtgt aggageagaa 360 acaaatggaa gtaaggatge gttgaataag tttteagata ataecetaaa getgttaaag 420

```
gagttaacaa gcaaaaaatc tcttcaagtg ccaagtattt attatcattt gcctcattta 480
ttgaaaaatg aaggaagtct tcaacctgct gtacagattg gcaacggaag aacaggagtt 540
tcaatagtca tgggcattcc cacagtgaag agagaagtta aatcttacct catagaaact 600
cttcattccc ttattgataa cctgtatcct gaagagaagt tggactgtgt tatagtagtc 660
ttcataggag agacagatat tgattatgta catggtgttg tagccaacct ggagaaagaa 720
ttttctaaag aaatcagttc tggcttggtg gaagtcatat cacccctga aagctattat 780
cctgacttga caaacctaaa ggagacattt ggagactcca aagaaagagt aagatggaga 840
acaaagcaaa acctagatta ctgttttcta atgatgtatg ctcaagaaaa gggcatatat 900
tacattcagc ttgaagatga tattattgtc aaacaaaatt attttaatac cataaaaaat 960
tttgcacttc aactttcttc tgaggaatgg atgattctag agttttccca gctgggcttc 1020
attggtaaaa tgtttcaagc gccggatctt actctgattg tagaattcat attcatgttt 1080
tacaaggaga aacccattga ttggctcctg gaccatattc tctgggtgaa agtctgcaac 1140
cctgaaaaag atgcaaaaca ttgtgataga cagaaagcaa atctgcgaat tcgcttcaga 1200
cettecettt tecaacatgt tggtetgeae teateactat caggaaaaat ceaaaaacte 1260
acggataaag attatatgaa accattactt cttaaaatcc atgtaaaccc acctgcggag 1320
gtatctactt cettgaaggt etaceaaggg cataegetgg agaaaaetta catgggagag 1380
gatttettet gggetateae acegataget ggagaetaea tettgtttaa atttgataaa 1440
ccagtcaatg tagaaagtta tttgttccat agcggcaacc aagaacatcc tggagatatt 1500
ctgctaaaca caactgtgga agttttgcct tttaagagtg aaggtttgga aataagcaaa 1560
gaaaccaaag acaaacgatt agaagatggc tatttcagaa taggaaaatt tgagaatggt 1620
gttgcagaag gaatggtgga tccaagtctc aatcccattt cagcctttcg actttcagtt 1680
attcagaatt ctgctgtttg ggccattctt aatgagattc atattaaaaa agccaccaac 1740
tgatcatctg agaaaccaac acattttttc ctgtgaattt gttaattaaa gatagttaag 1800
catgtatett ttttttattt etaettgaae actaeetett gtgaagteta etgtagataa 1860
gacgattgtc atttccactt ggaaagtgaa tctcccataa taattgtatt tgtttgaaac 1920
taagctgtcc tcagatttta acttgactca aacatttttc aattatgaca gcctgttaat 1980
atgacttgta ctattttggt attatactaa tacataagag ttgtacatat tgttacattc 2040
tttaaatttg agaaaaacta atgttacata cattttatga agggggtact tttgaggttc 2100
acttatttta ctatt
                                                                   2115
<210> 99
<211> 535
<212> PRT
<213> Homo sapiens
<400> 99
Met Arg Leu Arg Asn Gly Thr Val Ala Thr Ala Leu Ala Phe Ile Thr
                                    10
Ser Phe Leu Thr Leu Ser Trp Tyr Thr Thr Trp Gln Asn Gly Lys Glu
                                25
Lys Leu Ile Ala Tyr Gln Arg Glu Phe Leu Ala Leu Lys Glu Arg Leu
                            40
Arg Ile Ala Glu His Arg Ile Ser Gln Arg Ser Ser Glu Leu Asn Thr
                        55
Ile Val Gln Gln Phe Lys Arg Val Gly Ala Glu Thr Asn Gly Ser Lys
                   70
Asp Ala Leu Asn Lys Phe Ser Asp Asn Thr Leu Lys Leu Leu Lys Glu
```

```
Ile Gly Glu Thr Asp Ile Asp Tyr Val His Gly Val Val Ala Asn Leu
                                185
Glu Lys Glu Phe Ser Lys Glu Ile Ser Ser Gly Leu Val Glu Val Ile
                            200
Ser Pro Pro Glu Ser Tyr Tyr Pro Asp Leu Thr Asn Leu Lys Glu Thr
                        215
Phe Gly Asp Ser Lys Glu Arg Val Arg Trp Arg Thr Lys Gln Asn Leu
                    230
                                         235
Asp Tyr Cys Phe Leu Met Met Tyr Ala Gln Glu Lys Gly Ile Tyr Tyr
                245
                                     250
Ile Gln Leu Glu Asp Asp Ile Ile Val Lys Gln Asn Tyr Phe Asn Thr
                                265
Ile Lys Asn Phe Ala Leu Gln Leu Ser Ser Glu Glu Trp Met Ile Leu
                            280
Glu Phe Ser Gln Leu Gly Phe Ile Gly Lys Met Phe Gln Ala Pro Asp
                        295
                                             300
Leu Thr Leu Ile Val Glu Phe Ile Phe Met Phe Tyr Lys Glu Lys Pro
                    310
                                         315
Ile Asp Trp Leu Leu Asp His Ile Leu Trp Val Lys Val Cys Asn Pro
                325
                                    330
Glu Lys Asp Ala Lys His Cys Asp Arg Gln Lys Ala Asn Leu Arg Ile
            340
                                345
                                                     350
Arg Phe Arg Pro Ser Leu Phe Gln His Val Gly Leu His Ser Ser Leu
        355
                            360
                                                 365
Ser Gly Lys Ile Gln Lys Leu Thr Asp Lys Asp Tyr Met Lys Pro Leu
                        375
                                            380
Leu Leu Lys Ile His Val Asn Pro Pro Ala Glu Val Ser Thr Ser Leu
                    390
                                        395
Lys Val Tyr Gln Gly His Thr Leu Glu Lys Thr Tyr Met Gly Glu Asp
                405
                                    410
Phe Phe Trp Ala Ile Thr Pro Ile Ala Gly Asp Tyr Ile Leu Phe Lys
            420
                                425
Phe Asp Lys Pro Val Asn Val Glu Ser Tyr Leu Phe His Ser Gly Asn
                            440
                                                 445
Gln Glu His Pro Gly Asp Ile Leu Leu Asn Thr Thr Val Glu Val Leu
                        455
                                            460
Pro Phe Lys Ser Glu Gly Leu Glu Ile Ser Lys Glu Thr Lys Asp Lys
                    470
                                        475
Arg Leu Glu Asp Gly Tyr Phe Arg Ile Gly Lys Phe Glu Asn Gly Val
                485
                                    490
                                                         495
Ala Glu Gly Met Val Asp Pro Ser Leu Asn Pro Ile Ser Ala Phe Arg
            500
                                505
                                                    510
Leu Ser Val Ile Gln Asn Ser Ala Val Trp Ala Ile Leu Asn Glu Ile
                            520
                                                 525
His Ile Lys Lys Ala Thr Asn
                        535
```

```
<210> 100
```

<400> 100

```
attgctagag agagatggct ttcttttctc cctggaagtt gtcctctcag aagctgggct 60 ttttcctggt gactttcggc ttcatctggg gcatgatgct tctgcacttc accatccagc 120 agcggactca gcccgagagc agctccatgt tacgggagca gatccttgac ctcagcaaga 180 ggtacattaa ggcactggca gaggagaaca gggacgtggt ggatggcccc tacgctggtg 240
```

<211> 3226

<212> DNA

<213> Mus musculus

```
teatgacage etatgatetg aagaaaaege tegeegtett getggataae ateetgeage 300
gcattggcaa gctcgagtca aaggtggaca atctggtcaa cggcacagga gcgaactcca 360
ccaactccac cacggctgtc cccagcttgg tgtcgcttga gaaaattaat gtggcagata 420
tcattaatgg agttcaggaa aaatgtgtat tgcctcctat ggatggctac ccccactgcg 480
aggggaaaat caagtggatg aaggacatgt ggcgctcgga cccctgctac gcagactatg 540
gagtggacgg gacctcctgc tcctttttta tttacctcag tgaggttgaa aattggtgtc 600
ctcgtttacc ttggagagca aaaaatccct atgaagaagc tgatcataac tcattggcgg 660
aaatccgtac ggattttaac attctctacg gcatgatgaa gaagcacgag gagttccgtt 720
ggatgagget teggateegg egaatggetg aegegtggat ceaagetate aagtetetgg 780
cggagaaaca aaaccttgag aagaggaaac ggaagaaaat ccttgttcac ctggggctcc 840
tgaccaagga atcgggcttc aagattgcgg agacagcatt cagcggtggc cctctgggtg 900
aactcgttca gtggagtgac ttaatcacat ctctgtacct gctgggccat gacatccgga 960
teteggeete aetggetgag eteaaggaga taatgaagaa ggttgttgga aaceggtetg 1020
gctgtccaac tgtaggagac agaatcgttg agctgattta tatcgatatt gtgggacttg 1080
ctcaatttaa gaaaacacta gggccatcct gggttcatta ccagtgcatg ctccgggtgc 1140
tagactcctt tggaacagaa cctgagttca atcatgcgag ctatgcccag tcaaaaggcc 1200
acaagacccc ctggggaaag tggaatctga acccgcagca gttttacacc atgttccctc 1260
ataccccaga caacagcttt ctgggcttcg tggtggagca gcacctgaac tccagcgaca 1320
ttcaccacat caacgagatc aaaaggcaga accagtccct tgtgtatggc aaagtggata 1380
gtttctggaa gaataagaaa atctacctgg atatcattca cacgtacatg gaagtgcacg 1440
ccactgttta tggctccagt accaagaaca ttcccagtta cgtgaaaaac catggcattc 1500
tcagtggacg tgacctgcag tttcttctcc gggaaaccaa gctgttcgtt gggctcggat 1560
tecettatga aggeceaget eeeetggagg eeategegaa tggatgtget tteetgaace 1620
ccaagttcaa ccctcccaaa agcagcaaaa acacagactt cttcattggc aagccaacac 1680
tgagagaget gacateceag cateettaeg cagaagtett categgeegg ceacaegtet 1740
ggactgtgga teteaataac egagaggaag tagaagatge agtaaaagee atettaaace 1800
agaagattga gccgtatatg ccatatgagt tcacatgtga aggcatgctg cagagaatca 1860
acgettteat tgaaaaacag gaettetgee atggeeaagt gatgtggeeg eeceteageg 1920
ccctgcaggt taagctggct gagccagggc agtcctgcaa acaggtgtgc caggagagcc 1980
ageteatetg egageeatee ttettteaae aceteaaeaa qqaaaaqqae etqetqaagt 2040
ataaggtgac ctgccaaagc tcagaactgt acaaggacat cctggtgccc tccttctacc 2100
ccaagagcaa gcactgtgtg ttccaagggg acctcctgct cttcagttgt gccggagccc 2160
atcccacaca ccageggate tgcccetgee gggaetteat caagggeeaa gtggeeetet 2220
gcaaagactg cetatagcat egetgeeetg aattaactea gaegggaaag aegtggetee 2280
actgggcagg gccaaggggc acaaagacat tcagggactc tgaccagagc ctgagatctt 2340
tggtccaggg cttgagttta gtaccgctcc agccacagcc agtgcatccc agtttacacc 2400
aaaaccacaa gggaacaggt tagaacagga acctgggttc tcctcagtgt aaggaatgtc 2460
ctctctgtct gggagatcga gcgactgtag ggaaaggatc caggcagttg ctcccgggaa 2520
ttttttttt tttttttt aaagaaggga taaaagtccg gagactcatt caaactgaaa 2580
acaaaacagg aagagggaat tgagccaatt gggaaggact ttggggccga tcctaaacca 2640
attaatttat ttatttggga ggatggggc gggctcggga gggaggagag gggttgaaca 2700
gtttcctttt gttcctcact gttaattcgc ccaccttcgg gcccttcttg ttctgcagcg 2760
ccaagcaggg tgcagagggg ctgtggcttg cttgaggggc cactgtgggg cttcactcct 2820
ggtcacaggt ggcagcagag aaaagagatg tctataagca gggggatgta gctcagtttg 2880
tagaatgett geatageata aatgaagtee tgggtteeat eeceageace acataaatge 2940
aggtaagaaa cagagtcagg aggaccaagc attctccttg gctacataac aaaagcaagg 3000
cctttgtccc catgtcttgg ctacaagaga ccctatctca gaaaattgtg ggggggaggg 3060
ggggggaaat ggccttgaaa acacagccag tcactgtcac tgcattgcca gaactggtgg 3120
atcccaggtg tgcttggcag ataacagcta aaaggcacat aaccttggtg gggaaataaa 3180
tgcctgtggt gtcctgaggg ccccaccaag ttccaaaaaa aaaaaa
                                                                  3226
<210> 101
<211> 740
```

<212> PRT

<213> Mus musculus

<400> 101

Met Ala Phe Phe Ser Pro Trp Lys Leu Ser Ser Gln Lys Leu Gly Phe

```
Phe Leu Val Thr Phe Gly Phe Ile Trp Gly Met Met Leu Leu His Phe
                                25
Thr Ile Gln Gln Arg Thr Gln Pro Glu Ser Ser Ser Met Leu Arg Glu
Gln Ile Leu Asp Leu Ser Lys Arg Tyr Ile Lys Ala Leu Ala Glu Glu
                        55
Asn Arg Asp Val Val Asp Gly Pro Tyr Ala Gly Val Met Thr Ala Tyr
                                        75
                    70
Asp Leu Lys Lys Thr Leu Ala Val Leu Leu Asp Asn Ile Leu Gln Arg
Ile Gly Lys Leu Glu Ser Lys Val Asp Asn Leu Val Asn Gly Thr Gly
                                105
Ala Asn Ser Thr Asn Ser Thr Thr Ala Val Pro Ser Leu Val Ser Leu
                            120
Glu Lys Ile Asn Val Ala Asp Ile Ile Asn Gly Val Gln Glu Lys Cys
                        135
Val Leu Pro Pro Met Asp Gly Tyr Pro His Cys Glu Gly Lys Ile Lys
                   150
                                        155
Trp Met Lys Asp Met Trp Arg Ser Asp Pro Cys Tyr Ala Asp Tyr Gly
                                   170
                165
                                                        175
Val Asp Gly Thr Ser Cys Ser Phe Phe Ile Tyr Leu Ser Glu Val Glu
           180
                               185
Asn Trp Cys Pro Arg Leu Pro Trp Arg Ala Lys Asn Pro Tyr Glu Glu
       195
                           200
                                                205
Ala Asp His Asn Ser Leu Ala Glu Ile Arg Thr Asp Phe Asn Ile Leu
                       215
                                            220
Tyr Gly Met Met Lys Lys His Glu Glu Phe Arg Trp Met Arg Leu Arg
                   230
                                        235
Ile Arg Arg Met Ala Asp Ala Trp Ile Gln Ala Ile Lys Ser Leu Ala
                                    250
Glu Lys Gln Asn Leu Glu Lys Arg Lys Arg Lys Ile Leu Val His
                               265
Leu Gly Leu Leu Thr Lys Glu Ser Gly Phe Lys Ile Ala Glu Thr Ala
                           280
Phe Ser Gly Gly Pro Leu Gly Glu Leu Val Gln Trp Ser Asp Leu Ile
                        295
                                            300
Thr Ser Leu Tyr Leu Leu Gly His Asp Ile Arg Ile Ser Ala Ser Leu
                   310
                                        315
Ala Glu Leu Lys Glu Ile Met Lys Lys Val Val Gly Asn Arg Ser Gly
                325
                                    330
Cys Pro Thr Val Gly Asp Arg Ile Val Glu Leu Ile Tyr Ile Asp Ile
           340
                                345
Val Gly Leu Ala Gln Phe Lys Lys Thr Leu Gly Pro Ser Trp Val His
                            360
Tyr Gln Cys Met Leu Arg Val Leu Asp Ser Phe Gly Thr Glu Pro Glu
                        375
                                            380
Phe Asn His Ala Ser Tyr Ala Gln Ser Lys Gly His Lys Thr Pro Trp
                   390
                                        395
Gly Lys Trp Asn Leu Asn Pro Gln Gln Phe Tyr Thr Met Phe Pro His
               405
                                    410
Thr Pro Asp Asn Ser Phe Leu Gly Phe Val Val Glu Gln His Leu Asn
                                                    430
           420
                               425
Ser Ser Asp Ile His His Ile Asn Glu Ile Lys Arg Gln Asn Gln Ser
                           440
                                                445
Leu Val Tyr Gly Lys Val Asp Ser Phe Trp Lys Asn Lys Lys Ile Tyr
                       455
                                            460
Leu Asp Ile Ile His Thr Tyr Met Glu Val His Ala Thr Val Tyr Gly
```

```
470
                                        475
Ser Ser Thr Lys Asn Ile Pro Ser Tyr Val Lys Asn His Gly Ile Leu
                485
                                    490
Ser Gly Arg Asp Leu Gln Phe Leu Leu Arg Glu Thr Lys Leu Phe Val
                                 505
Gly Leu Gly Phe Pro Tyr Glu Gly Pro Ala Pro Leu Glu Ala Ile Ala
                            520
Asn Gly Cys Ala Phe Leu Asn Pro Lys Phe Asn Pro Pro Lys Ser Ser
                        535
                                             540
Lys Asn Thr Asp Phe Phe Ile Gly Lys Pro Thr Leu Arg Glu Leu Thr
                    550
                                         555
Ser Gln His Pro Tyr Ala Glu Val Phe Ile Gly Arg Pro His Val Trp
                565
                                     570
Thr Val Asp Leu Asn Asn Arg Glu Glu Val Glu Asp Ala Val Lys Ala
                                 585
Ile Leu Asn Gln Lys Ile Glu Pro Tyr Met Pro Tyr Glu Phe Thr Cys
                             600
Glu Gly Met Leu Gln Arg Ile Asn Ala Phe Ile Glu Lys Gln Asp Phe
                        615
                                             620
Cys His Gly Gln Val Met Trp Pro Pro Leu Ser Ala Leu Gln Val Lys
                    630
                                         635
Leu Ala Glu Pro Gly Gln Ser Cys Lys Gln Val Cys Gln Glu Ser Gln
                645
                                     650
Leu Ile Cys Glu Pro Ser Phe Phe Gln His Leu Asn Lys Glu Lys Asp
                                665
Leu Leu Lys Tyr Lys Val Thr Cys Gln Ser Ser Glu Leu Tyr Lys Asp
        675
                            680
                                                 685
Ile Leu Val Pro Ser Phe Tyr Pro Lys Ser Lys His Cys Val Phe Gln
                        695
                                            700
Gly Asp Leu Leu Phe Ser Cys Ala Gly Ala His Pro Thr His Gln
                   710
                                        715
Arg Ile Cys Pro Cys Arg Asp Phe Ile Lys Gly Gln Val Ala Leu Cys
Lys Asp Cys Leu
<210> 102
<211> 4
<212> PRT
<213> Artificial Sequence
<220>
<223> Illustrative retention signal peptide
<400> 102
Lys Asp Glu Leu
<210> 103
<211> 60
<212> PRT
<213> Saccharomyces cerevisiae
<400> 103
Ile Pro Phe Val Leu Ile Ala Ser Asn Phe Ile Gly Val Leu Phe Ser
```

```
Arg Ser Leu His Tyr Gln Phe Leu Ser Trp Tyr His Trp Thr Leu Pro
                                 25
Ile Leu Ile Phe Trp Ser Gly Met Pro Phe Phe Val Gly Pro Ile Trp
                             40
Tyr Val Leu His Glu Trp Cys Trp Asn Ser Tyr Pro
                         55
<210> 104
<211> 58
<212> PRT
<213> Drosophila virilis
<400> 104
Leu Pro Phe Phe Leu Cys Asn Phe Ile Gly Val Ala Cys Ala Arg Ser
                                     10
Leu His Tyr Gln Phe Tyr Ile Trp Tyr Phe His Ser Leu Pro Tyr Leu
                                 25
Val Trp Ser Thr Pro Tyr Ser Leu Gly Val Arg Tyr Leu Ile Leu Gly
                             40
Ile Ile Glu Tyr Cys Trp Asn Thr Tyr Pro
                        55
<210> 105
<211> 60
<212> PRT
<213> Saccharomyces cerevisiae
<400> 105
Ile Pro Phe Val Leu Ile Ala Ser Asn Phe Ile Gly Val Leu Phe Ser
Arg Ser Leu His Tyr Gln Phe Leu Ser Trp Tyr His Trp Thr Leu Pro
Ile Leu Ile Phe Trp Ser Gly Met Pro Phe Phe Val Gly Pro Ile Trp
                            40
Tyr Val Leu His Glu Trp Cys Trp Asn Ser Tyr Pro
<210> 106
<211> 59
<212> PRT
<213> Drosophila melanogaster
<400> 106
Leu Pro Phe Phe Leu Cys Asn Leu Val Gly Val Ala Cys Ala Ser Arg
Ser Leu His Tyr Gln Phe Tyr Val Trp Tyr Phe His Ser Leu Pro Tyr
            20
                                25
Leu Ala Trp Ser Thr Pro Tyr Ser Leu Gly Val Arg Cys Leu Ile Leu
                            40
Gly Leu Ile Glu Tyr Cys Trp Asn Thr Tyr Pro
```